

HYDROXYLASES AND MODULATORS THEREOF

Field of Invention

The present invention relates to methods of designing inhibitors of FIH using
5 the crystal structure of FIH, and to inhibitors of FIH and their use in the treatment of
ischaemia.

Background of the Invention

In cells of many organisms exposure to an environment in which oxygen is
10 depleted relative to optimal levels induces a hypoxic response. In these hypoxic
cells, activation of a transcriptional cascade involving hypoxia inducible factor (HIF)
directs a series of adaptive responses that enhance oxygen delivery or limit oxygen
demand. Activation of HIF in cancer and ischaemic hypoxic vascular diseases has
revealed its important role in human pathology and demonstrated that manipulation
15 of HIF activity has important therapeutic potential.

The HIF transcriptional complex comprises an $\alpha\beta$ heterodimer, HIF- β being a
constitutive nuclear protein that dimerises with oxygen regulated HIF- α subunits
(Semenza, G. L. (2000) *Genes Dev.* 14, 19831991). The activity of HIF- α , is
suppressed by oxygen-dependent modification catalysed by a series of Fe^(II) and 2OG
20 dependent dioxygenases that hydroxylate specific HIF- α residues. In the presence of
oxygen in human HIF-1 α , 4-hydroxylation of Pro402 or Pro564 by a set of HIF
prolyl hydroxylase isozymes (PHD1-3) (Epstein et al. (2001) *Cell* 107, 4354; Bruick,
R. K., and McKnight, S. L. (2001) *Science* 294, 13371340) mediates its recognition
by the von Hippel-Lindau (VHL) ubiquitin ligase complex and consequent targeting
25 for proteasomal destruction (Ivan et al, (2001) *Science* 292, 464468; Jaakkola et al
(2001) *Science* 292, 468472, WO 02/074981). In a complementary mechanism FIH
catalyses β -hydroxylation of HIF-1 α Asn803 (Lando et al, (2002) *Science* 295,
858861) blocking interaction with the transcriptional co-activator p300 (Dames et al.,
(2002) *Proc. Natl. Acad. Sci. U. S. A.* 99, 52715276; Freedman et al, (2002) *Proc.*
30 *Natl. Acad. Sci. U. S. A.* 99, 53675372). In hypoxia, limitation of enzymatic activity
allows HIF- α to escape destruction and become transcriptionally active.

Inhibition of HIF hydroxylases strongly activates the HIF transcriptional cascade even in the presence of oxygen (Epstein et al.(2001) *Cell* 107, 4354). Thus, inhibition of the HIF hydroxylases results in a pro-angiogenic response that may be used in the treatment of cardiovascular diseases/ ischaemic hypoxic vascular diseases including myocardial infarction and anaemia. A problem with this approach is that the human cells contain other enzymes belonging to the same family as the HIF hydroxylases, i.e. utilising dioxygen (a cosubstrate), 2-oxoglutarate (2OG) (a cosubstrate) and Fe(II) (a cofactor). Such enzymes are exemplified by phytanoyl coenzyme A hydroxylase, procollagen prolyl-4-hydroxylase, procollagen prolyl-3-hydroxylase, gamma-butyrobetaine hydroxylase, Alk B (a DNA repair enzyme) and others including predicted 2OG oxygenases identified on the basis of sequence analyses including a sub-family related to FIH (Hewitson et al., *J BIOL CHEM* 277 (29): 26351-26355, 2002). It is generally agreed that it is desirable that enzyme inhibitors used as pharmaceuticals are selective for their intended target or the targets involved in producing the desired effect. A lack of selectivity can lead to toxic side effects that render particular compounds unsuitable for use in human or animal therapy. One approach to identifying compounds that are selective for the intended target is to undertake structural, mechanistic and other analyses on the intended agents and to use the information gained to aid in the preparation of selective compounds, or more selective compounds (relative to those previously known), for use as pharmaceuticals for use in humans or animals. Here we describe structural and other studies on the HIF hydroxylases that enable the design of selective inhibitors of FIH and related enzymes.

Summary of the Invention

The present inventors have now identified the site of hydroxylation of asparagine 803 of HIF-1 α by FIH. In addition, the inventors have obtained the crystal structure for FIH including identification of the binding site and residues involved in the interaction of FIH with HIF.

Accordingly, the present invention provides a method of identifying, screening, characterising or designing a chemical entity which mimics or binds to FIH, which method comprises comparing a structural model of FIH with a structural

model for said chemical entity, wherein said structural model of FIH is derived from structural factors or structural coordinates determined by subjecting to X-ray diffraction measurements a crystal comprising FIH.

The invention also provides for:

- 5 - the use of the structural co-ordinates obtainable by subjecting a crystal comprising FIH to X-ray diffraction measurements and deducing the structural co-ordinates from the diffraction measurements, to identify, screen, characterise, design or modify a chemical entity;
- a chemical entity identified by a method of the invention, wherein
- 10 said chemical entity inhibits the asparaginyl hydroxylase activity of FIH; and
- a chemical entity of the invention for use in a method of treatment.

Description of the Figures

Figure 1: 2OG binding site.

15 Figure 2: binding of Asn-803.

Figure 3: conformation of CAD at site 1.

Figure 4: conformation of CAD at site 2.

Figure 5: figure indicating the turn formed by 802-804 of HIF-CAD at the active site of FIH.

20 Figure 6: conformation of the turn formed by residues 802-804 of HIF-CAD at the active site of FIH.

Detailed Description of the Invention

The present inventors have identified the position of asparagine 803 that is

25 hydroxylated by FIH. In addition, the inventors have identified the crystal structure of FIH. This structure therefore allows for identification of the amino acid residues involved in binding of FIH to HIF.

The identification of the interaction and the structures allows for the characterisation or identification of chemical entities which can bind and in particular

30 which can inhibit FIH. A number of different types of inhibitors can be identified as discussed in more detail below.

The inventors have successfully crystallised human FIH. This the first crystallisation of FIH and has enabled determination of the crystal structure. Co-ordinates from the crystal analysis are set out in Table 3 below. The studies have allowed analysis of the binding of asparagine-803 of HIF and analysis of the conformation of the c-terminal activation domain (CAD) of HIF at the binding sites to FIH. The present invention provides the use of the structural co-ordinates of FIH to identify, characterise, design or screen chemical entities. The chemical entities of interest are those which bind to FIH and in particular which inhibit the asparaginyl hydroxylase activity of FIH. In addition, chemical entities may be identified, characterised or designed which are modified asparagine hydroxylases.

Typically, the structural co-ordinates used are obtainable by subjecting a crystal comprising FIH or a fragment thereof to X-ray diffraction measurements and deducing the structural co-ordinates from the diffraction measurements, to identify, screen, characterise, design or modify a chemical entity. The structural co-ordinates indicate the positions of individual atoms within the crystal and give an indication of the space available for adjusting the position of individual atoms when designing a chemical entity.

The crystal subjected to X-ray diffraction methods comprises FIH or a fragment thereof. The FIH may be from any source but is preferably human FIH. The FIH may be a modified form. For example, the FIH may be modified by insertion, deletion, n-terminal or C-terminal addition, or substitution of amino acid by another amino acid. Amino acid substitutions may be conservative substitutions. Typically, when crystallised, a FIH mutant will adopt a similar 3-dimensional structure to that adopted by the corresponding FIH. A mutant may be an inactive FIH.

References to FIH herein refer to FIH and homologues thereof. Amino acid residues are defined with reference to the position in FIH (see e.g. Hewitson et al). The relevant amino acid residues of homologues of FIH are the equivalent amino acid residues, based on for example the best alignment of homologue to FIH.

A FIH may be isolated by any suitable means for use in crystallisation studies. For example, a FIH may be purified using biochemical means from a suitable source. Typically, however, it will be convenient to over express FIH in cells and purify FIH from those cells. Thus, a polynucleotide encoding a FIH may be used

in the construction of a vector. The FIH may be crystallised according to any method known to those skilled in the art. X-ray diffraction may be carried according to any suitable method. The data collected from X-ray diffraction experiments may be processed to deduce the structural co-ordinates of FIH using any suitable method.

5 The invention provides the use of structural co-ordinates to identify, characterise, design or screen a chemical entity. The chemical entity may be one which binds to FIH, or which acts as an inhibitor of asparaginyl hydroxylase activity. Alternatively, the chemical entity may be a modified FIH to alter the activity of a FIH.

10 A chemical entity which binds to or inhibits FIH is any chemical entity capable of forming an association with the FIH. The binding or inhibition may be non-specific, for example, such an entity may also bind to or inhibit other 2OG oxygenases. Alternatively, an agent may be designed or identified which specifically binds to or inhibits asparaginyl hydroxylases. An agent may be designed or identified
15 which is a specific inhibitor of FIH, but not other asparaginyl hydroxylases.

 The structural co-ordinates of FIH allows a skilled person to predict which amino acids are important in active site formation and which amino acids are important in contacting the substrate. The substrate binding site may be shown as a 2 dimensional representation or a 3 dimensional representation produced by physical
20 models or displayed on a computer screen. Such representations can be used to design, identify or screen chemical entities which bind to or inhibit or are predicted to bind to or inhibit FIH. Such representations can also be used to identify modifications of FIH to alter its activity characteristics.

 Examples of modifications to FIH include modifications to increase the
25 binding of FIH for its substrate, or to alter the substrate the specificity. Alternative modifications include those which alter the activity of FIH, for example, to remove asparaginyl hydroxylase activity.

 The representations of the structures may be used in other ways. For example, the representations of the FIH active site may be used to model constraints
30 by the putative introduction of covalent bonds between the atoms which come close together when FIH binds to a substrate. Representation of the active site may be used to predict which residues of FIH are likely to be involved in steric hindrance.

Such residues may be modified, replaced or deleted to decrease esoteric hindrance in order to increase avidity of the peptide for its substrates.

In general, it will be necessary to process the structural co-ordinates obtainable according to the invention in computer-based methods in order to identify or design chemical entities with the desired molecular structure or to identify
5 chemical entities whose structure is complementary to all or part of another chemical entity of interest. Thus, chemical entities which have a structure similar to FIH may be identified or designed. Chemical entities which bind to FIH may be identified or designed. Preferably, such chemical entities bind at the active site of FIH and in
10 general may act as inhibitors of asparaginyl hydroxylase activity.

Such computer-based methods fall into two broad classes: database methods and *de novo* designed methods. In database methods, the chemical entity of interest is compared to all chemical entities present in a database of chemical structures and chemical identities whose structure is in some way similar to the compound of
15 interest identified. The structures in the database are based either on experimental data, generated by NMR or X-ray crystallography, or models of 3 dimensional structures based on 2 dimensional data. In *de novo* design methods, models of chemical entities, for example such are those which might bind to FIH are generated by a computer program using information derived from known structures and/or
20 theoretical rules.

Similarly, the FIH structural coordinates may be used to screen for the expected activity of chemical entities selected, designed or shown to be modulators such as inhibitors of other hydroxylases, for example prolyl hydroxylases. For example the compounds may be screened to assess the likelihood of a prolyl
25 hydroxylase inhibitor additionally inhibiting FIH hydroxylase. Such screening methods may be useful in identifying agents which selectively inhibit HIF prolyl hydroxylase, but not HIF asparaginyl hydroxylase.

Chemical entities designed or selected according to the methods of the invention may be tested and optimised using computational or experimental
30 evaluation. Experimental methods to assay for the activity of asparaginyl hydroxylase are described in more detail below.

Based on the structure of FIH, a number of different types of inhibitors can be identified. These inhibitors are discussed in more detail below.

Dimerisation inhibitors

5 The crystallographic asymmetric unit contains one FIH molecule. However, analysis of crystallographic symmetry revealed a dimeric form of FIH, consistent with native gel-electrophoresis analysis. The dimer interface involves the two C-terminal helices of each molecule in an interlocking arrangement predominantly involving hydrophobic interactions. This unusual interface buries a surface area of
10 3210 Å², large on average by comparison to other dimeric proteins of this size. Inhibitors of dimerisation include those that bind to residues that form the dimerisation interface including residues selected from 330-346, such as Leu-340 and Ile-344. Inhibitors include peptides or peptide mimetics that correspond to all or part of the FIH residues involved in the dimerisation interface.

15 For example, such inhibitors may comprise a fragment of FIH, for example, including the residues from 340 to 344, preferably, including residues 330 to 346. Such a fragment may typically have 6 or 10 amino acids in length, preferably, up to 15 or 20 amino acids in length. Alternatively, peptide homologues may be used, for example, which comprise a homologue to the residues of 340 to 344 or 330 to 336,
20 including 1, 2 or more substitutions. Additional agents include peptides or peptide mimetics which can be designed based on the crystal structure to interfere with dimerisation.

Inhibitors exploiting metal binding in FIH:

25 The structural work defines the presence of Fe(II) at the active site of FIH and by implication related HIF hydroxylases. The iron is bound in an almost octahedral manner by the side chains of His199, Asp201 and His279, the 2-oxo and 1-carboxylate groups of 2OG. In the enzyme-substrate complexes there is a vacant position opposite His279 revealing that the enzyme is primed for dioxygen binding.
30 Accommodation of a ligand opposite His279 may require disruption of the hydrogen bond between Asp201 and CAD Asn803 (the iron and Asn803 β-carbon are only ~4.9 Å apart). Subsequent decarboxylation of 2OG presumably yields an iron-oxo

species $[\text{Fe}^{(\text{IV})}=\text{O} \leftrightarrow \text{Fe}^{(\text{III})}-\text{O}\cdot]$ that effects oxidation at the carbon of Asn-803 in the C-terminal transactivation domain (CAD) of HIF.

Compounds that contain functional groups that bind to iron are useful as inhibitors of FIH. Examples of such compounds include thiols, alcohols, phenols including flavonoids such as quercitin and derivatives thereof, carbohydrates, hydroxamates, imidazoles and other heterocycles for example nitrogen containing heterocycles.

$\text{Zn}^{(\text{II})}$ binds to FIH in an identical manner to $\text{Fe}^{(\text{II})}$ (structure 3), consistent with the metal-mediated hypoxic effect being due to displacement of $\text{Fe}^{(\text{II})}$ from the active site of HIF hydroxylases. Since neither $\text{Zn}^{(\text{II})}$ nor other metal inhibitors of FIH can replace $\text{Fe}^{(\text{II})}$ as a cofactor in catalysis, compounds that preferentially promote the binding of a metal other than iron [such as $\text{Zn}^{(\text{II})}$] at the active site of FIH act as inhibitors.

A further class of inhibitor are non-metallic inhibitors that operate via competing with $\text{Fe}^{(\text{II})}$ for binding at the active site. Such inhibitors may bind to any or all of the triad of residues (His-199, Asp-201, His-279), that bind the $\text{Fe}^{(\text{II})}$ at the active site of catalytically active FIH.

Inhibitors exploiting the 2OG binding sites

The FIH:CAD structures with NOG reveal that like 2OG it is ligated to iron in a bidentate manner and imply it is an inhibitor due to decreased susceptibility to attack by an iron bound (su)peroxide intermediate or by hindering binding of dioxygen to the metal.

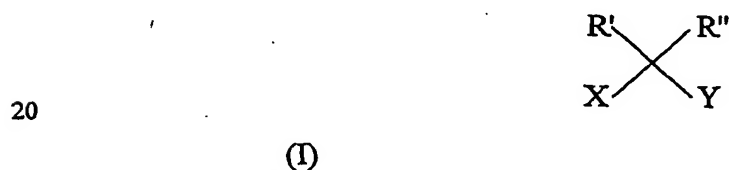
The structural studies on FIH reveal the binding interactions for the 2OG and NOG (see for example Figure 1). The 5-carboxylate of 2OG (and the equivalent carboxylate of NOG) forms hydrogen bonds with the side-chains of Lys214, Thr196 and Tyr145; such interactions are unprecedented in other structures of 2OG oxygenases. FIH is further unusual in that Lys214 is on the fourth DSBH (double stranded beta-helix) β -strand whereas previously assigned basic 2OG-5-carboxylate binding residues are at the beginning of the eighth DSBH strand.

The structural studies reveal the FIH residues that form the pocket into which 2OG and NOG bind. In addition to the aforementioned these include the side-chains -

of Ile-281, Leu-186, Leu-188, Phe-207, Thr-196. Knowledge of these interactions enables the design of improved (as measured by binding parameters) and selective inhibitors. Thus, for example an inhibitor binding in the 2OG binding pocket may form hydrophobic interactions with any or all of the side chains of Ile-281, Leu-186
 5 Leu-188, Phe-207, Thr-196. Further it may form electrostatic or hydrogen bonding interactions with the residues involved in binding the 5-carboxylate of 2OG (Lys214, Thr196 and Tyr145).

Selective inhibition of FIH via inhibitors interacting with the 2OG binding residues is exemplified as follows: kinetic analyses of a series of inhibitors based
 10 upon *N*-oxaloyl amino acids revealed the *R*-enantiomer (IC_{50} 0.4 mM) of *N*-oxaloylalanine was significantly more potent than the *S*-enantiomer (IC_{50} 2.5 mM). Analysis of the 2OG binding pocket in FIH reveals that the binding of the *S*-enantiomer is hindered by interactions between its methyl group and the side chain of Thr-196 and, Ile-281 in the 2OG binding pocket. A reversed selectivity (i.e. the *S*-
 15 enantiomer was more potent) was observed both for procollagen prolyl-hydroxylase and the PHD isozymes, demonstrating it should be possible to develop selective inhibitors for individual types of HIF hydroxylase. Such inhibitors may or may not chelate to an active site metal.

Compounds include those of general formula

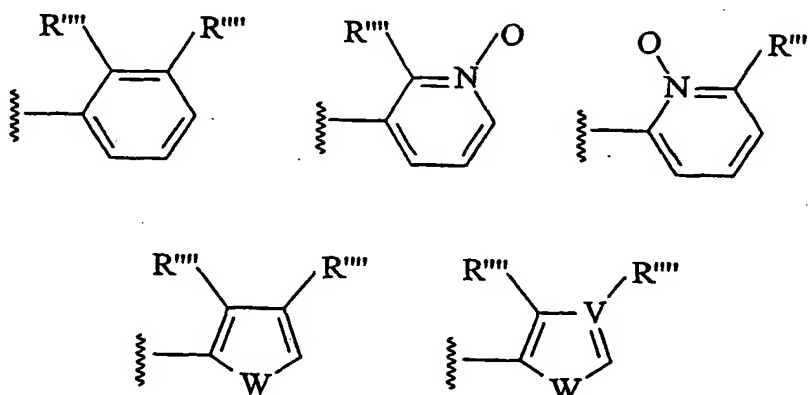


wherein each of R' and R'' , which may be the same or different, is H, F or C_1 to C_3 alkyl or substituted alkyl, CH_2OH , CH_2CO_2H or $CONH_2$, X is $COOH$, $SOOH$, or
 25 $CONHH$ or an ester thereof, or heterocyclic or other group which forms a favourable interaction with one or more of the side chains of Lys-214, Thr-196 and Tyr-145, i.e. those residues involved in binding the 5-carboxylate of 2OG as revealed in the crystallographic analyses,

Y is $-(CR'''R''')_nZ$, where Z is

-NR'''COCOOH, -NR'''CSCOOH, -NR'''COCOSH,
 -CHSR'''CONR'''R''', -CHOR'''CONR'''OR''', -CHSR'''CONR'''OR''' or
 -CHOR'''CONR'''NR'''OR''', wherein each R''', which may be the same or
 different, is H, alkyl, OH or O-alkyl, n is 0 to 3 and preferably 0, or

5



10 wherein R''' is OH, OR''' or NHCOR''', and W is S, NH, or O.

Thus X is a group that forms favourable interactions with one or more of the
 side chains of interactions one or more of the side chains of Lys-214, Thr-196 and
 Tyr-145, i.e. those residues involved in binding the 5-carboxylate of 2OG. X may be
 functionalised as a pro-drug such that is delivered to the desired site of action or has
 15 desirable pharmacokinetic properties. As indicated above, X can be an ester such a
 methyl or ethyl ester or amide derivative of carboxylic acid versions of X.

If n is 0, Y is typically CONHOH, CONHNH₂, NR'''COCOOH,
 NR'''CSCOOH or NR'''COCOSH. Y is preferentially of a size such that it can
 chelate to the active site metal whilst maintaining all or some of the favourable
 20 binding interactions found in the 2OG binding pocket as defined by crystallographic
 analyses. As with X, Y may be functionalised as a pro-drug.

When Y contains an aromatic ring as indicated above it can comprise other
 ring systems including aryl or functionalised aryl rings as well as heterocyclic and
 functionalised heterocyclic rings. The above rings may be further functionalised to
 25 optimise binding at the FIH active site.

Inhibitors exploiting the peptide substrate binding siteThere are two binding sites

The ES complex structures unexpectedly reveal two separate binding sites involving CAD₇₉₅₋₈₀₆ (i.e residues 795-806 of the C-terminal transactivation domain of HIF) (Site 1) and CAD₈₁₃₋₈₂₂ of HIF (Site 2) with contact surface areas of 1640 Å² and 1080 Å², respectively (see for example the figures). CAD residues in these regions are conserved in all known HIF-1α and HIF-2α sequences. The electron density for site 1 was of good quality, with only the side-chain of Tyr798 poorly defined, while that for site 2 was at a lower level and quality, probably reflecting weaker binding at this site. CAD₈₀₄₋₈₀₆ and presumably also CAD₈₀₇₋₈₁₁, for which density was not observed, do not form direct interactions with FIH. Kinetic analyses employed to investigate the relative importance of Sites 1 and 2, revealed that fragments containing site 1 only are hydroxylated by FIH but less efficiently than those containing both sites demonstrating that both are important in binding and that both may be exploited in inhibition studies.

At Site 1 CAD₇₉₅₋₈₀₃ are bound in a groove and adopt a largely extended conformation linked to FIH by ten hydrogen bonds. Asn803 of CAD is strikingly buried at the active and directly adjacent to the Fe^(III). CAD Asn803 and Ala804 form a tight turn, stabilised by a hydrogen bond between the backbone carbonyl of Val802 and NH of Ala804, which projects the side chain of Asn803 towards the Fe^(III). The side chain of CAD Asn803 is precisely orientated by three hydrogen bonds to enable hydroxylation at the *pro-S* position of the β-carbon consistent with the NMR assignments (see above) The primary amide of CAD Asn803 is sandwiched between FIH residue Tyr102 and the Fe^(III), and forms hydrogen bonds with the side chains of FIH residues Gln239 and Arg238, residues located on the insert to the DSBH motif. Significantly, the substrate and Fe^(III) binding sites are directly linked since the backbone nitrogen of CAD Asn803 also forms a hydrogen bond (~3 Å) with the carboxylate oxygen of Asp201 that is not complexed to the iron. Six additional hydrogen bonds stabilise the binding of FIH to CAD₇₉₅₋₈₀₁.

In contrast with Site 1, Site 2 is located on the FIH surface and involves only two hydrogen bonds. CAD₈₁₆₋₈₂₃ of Site 2 form an α-helix, in exact agreement with the structure of this region in complex with CBP/p300 (Dames et al., (2002) *Proc.*

Natl. Acad. Sci. U. S. A. **99**, 52715276; Freedman et al, (2002) *Proc. Natl. Acad. Sci. U. S. A.* **99**, 53675372). As in that complex, the highly conserved Leu818, Leu819 and Leu822 sit in a hydrophobic pocket on the surface of FIH and form the basis of the binding interaction and so it is not possible for these residues to bind
5 simultaneously to CBP/p300 and FIH.

The extended loop conformation adopted by the CAD residues at Site 1, contrasts with the α -helical conformation adopted by the same residues when complexed with the 1st transcriptional adaptor zinc-binding domain (TAZ1) of CBP/p300(Dames et al.,(2002) *Proc. Natl. Acad. Sci. U. S. A.* **99**, 52715276;
10 Freedman et al, (2002) *Proc. Natl. Acad. Sci. U. S. A.* **99**, 53675372). The disordered structure observed for the CAD, and other HIF- α residues, when free in solution may thus reflect a requirement to adopt more than one conformation for complex formation with different proteins.

The changes in the conformation of CAD on binding are complemented by
15 changes in FIH revealing an induced fit binding process; Trp296 of FIH undergoes a 50° rotation about C_{beta}-C_{alpha} to accommodate CAD Val802, while both Tyr102 and Tyr103 become more ordered. Further evidence of induced fit comes from the significant differences in resolution between the structures obtained with and without CAD fragments bound reflecting ordering of FIH that occurs on binding (structure 4,
20 for comparison, represents FIH complexed with Fe^(II) and 2OG alone). Interference in the conformational changes involved in the hypoxic response, in particular those involving the CAD region, e.g. by use of small molecules or by gene or protein therapy, may allow manipulation of the hypoxic response to enable pro or anti-angiogenic responses.

25 Thus, the structural studies define the (i) FIH residues involved in binding the CAD of HIF (ii) conformation of FIH when CAD is bound and (iii) conformation of CAD when bound to FIH. These results are useful in the design of selective inhibitors of FIH and related enzymes. Features of the FIH binding sites may be used to mediate tighter binding of inhibitors to FIH or to obtain inhibitors that do not bind
30 tightly to FIH, i.e. avoid inhibition of FIH.

Inhibitors binding at or close to the Site 1 may exploit electrostatic, hydrogen binding and/or hydrophobic interactions with Tyr-102, Asp-104, Lys-106, Asp-201,

Glu-202, Gln-147, Gln-239, residues 299-303, His-313, Ala-317, Ile-318, Asn-321, Lys-324, Arg-238, Trp-296, Asn-321- Lys-324. Inhibitors binding at Site 1 may mimic or partially mimic the turn conformation adopted by CAD when bound at Site 1.

5 Inhibitors binding at or close to Site 2 may exploit electrostatic, hydrogen binding and/or hydrophobic interactions with residues Thr-149, Leu-150, Asn- 151, Asp-152 and residues Val-159, Phe-162, Leu-163, Trp-167, Gln-181, Leu-182, Thr-183, Ser- 184, Asn- 185. Inhibitors binding at Site 2 may mimic or partially mimic the helical conformation adopted by CAD when bound at Site 2.

10 It is recognised that inhibitors need not bind to both Sites 1 and 2, although that they may, and that Site 1 is preferred over Site 2.

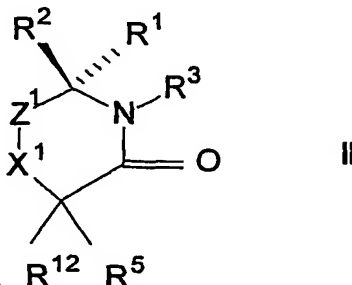
Residues 801 – 805 of CAD that bind at Site 1, and in particular residues 802-805 form a turn conformation in which the distance of the backbone C=O of 802 to the backbone NH of 804 is ca. 2.8 Å. Including the H-bond formed between the NH
15 of Ala-804 and the carbonyl O of Val-802 of the HIF-1 alpha CAD, the turn contains 7 atoms in a pseudo-ring.

Turns are especially amenable to mimicry by analogues useful for enzyme inhibition or receptor binding. The medicinal chemistry literature is replete with examples of such turn mimics. These can be modified by known methods to bind to
20 specific targets, in particular given the knowledge of the target structure.

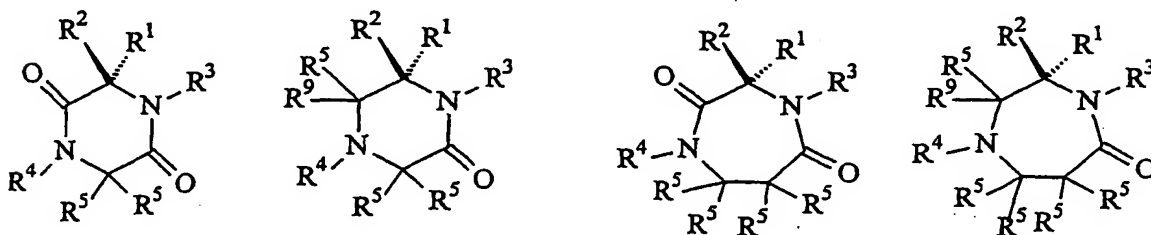
Examples of turn mimics and their modifications can be found in the following reviews: Hanessian et al, TETRAHEDRON 53: 12789-12854 SEP 22 1997; Gillespie et al, BIOPOLYMERS 43: 191-217 1997; and Burgess et al., ACCOUNTS CHEM RES 34: 826-835 2001). Recent examples of primary reports
25 on turns include the following (and references therein) Maier et al, EUR J ORG CHEM: 2686-2689, 2002; Reid et al, J AM CHEM SOC 124: 5673-5683, 2002; Mahadevan et al, J BIOMOL STRUCT DYN 19: 775-788 2002; Eguchi et al, J MED CHEM 45: 1395-1398 2002; De Borggraeve et al, TETRAHEDRON LETTERS 42: 5693-5695 2001; Kohn et al, TETRAHEDRON LETT 42: 4453-
30 4457 2001; Eguchi et al, TETRAHEDRON LETT 42: 1237-1239 2001; Manzoni et al, TETRAHEDRON 57: 249-255 2001; Jiang et al., HELV CHIM ACTA 83: 3097-3112 2000; Derrer et al, J CHEM SOC PERK T 1: 2957-2967 2000; Belvisi et al,

EUR J ORG CHEM: 2563-2569 2000; Claridge et al, BIOORG MED CHEM LETT
6: 485-490 1996.

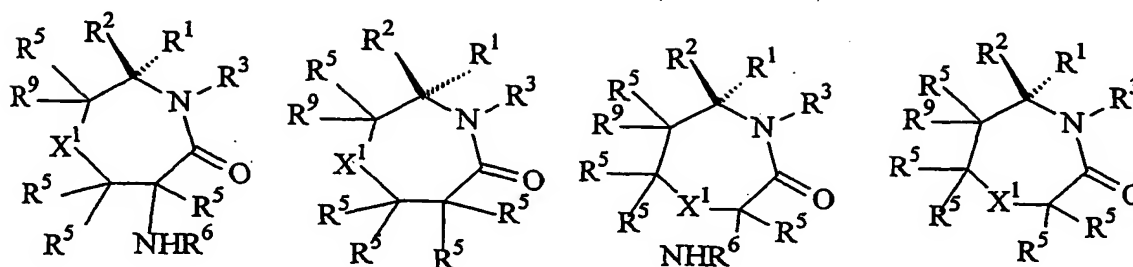
These include compounds of the general formula:



wherein R^1 is such that it can form an electrostatic or H-bonding interaction with Gln-237 and or Arg-238, preferably $CR^8R^9CONH_2$ or an analogue thereof where R^8 is hydrogen or a peptide or peptide mimetic (such as those composed of β -amino acids or peptide isosteres), and R^9 is hydrogen, optionally functionalised alkyl, optionally functionalised aryl, heteroaryl or any combination thereof such as CH_2CONH_2 , R^2 is hydrogen or a group that will interact favourably with Tyr-102 of FIH, R^3 is H or a group which can form a H-bond with Asp-201, Z^1 is $>C=O$ or $>CR^5R^9$ where R^5 is hydrogen, optionally functionalised alkyl, aryl, or heteroaryl or any combination thereof, R^{12} is as defined for R^5 or is NHR^6 where R^6 is COR^5 or SO_2R^5 and X^1 is NR^4 , $NR^4C(R^5)_2$, $C(R^5)_2NR^4$, or O or NH where R^4 is COR^5 or SO_2R^5 . In this and in the other formulae each $R^1, R^2, R^3, R^4, R^5, R^6, R^7, R^8, R^9, R^{10}, R^{11}$ and R^{12} can be the same or different. In particular, these compounds may have one of the formulae

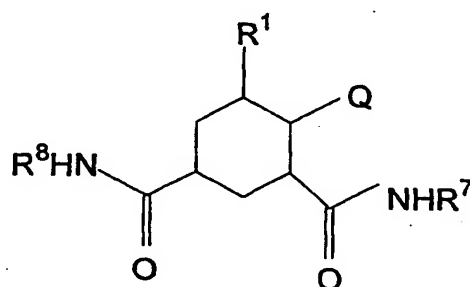


15



wherein the radicals are as defined above, and R^7 and R^8 are independently peptides or peptides mimetics or part peptide mimetics, such as those containing or consisting of beta-amino acid residues, urethane, sulphonamide or phosphonamide links.

Other compounds which can be used are those possessing the formula

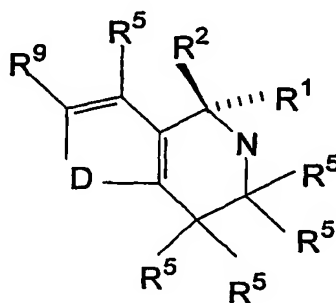


III

10

where Q represents H or OH and R^7 and R^8 are as defined above.

Further compounds which can be used possess the formula



IV

wherein R^1 , R^2 , R^5 and R^9 are as defined above and D is S, O, NH or $CHR^9=CHR^9$. Thus the ring attached to the six-membered ring is either a five-membered heterocyclic ring or an aryl ring.

In these formulae R^8 and R^9 can be optimised to bind in the channel linking the 2OG and peptide substrate binding sites and to the 2OG binding site itself.

Cyclic peptides acting as mimics of the turn adopted by CAD in site 1. The cyclo may be formed via peptide links, disulphide bonds or C-C bonds.

Inhibitors employing a combination of binding sites

It is well known that enzyme inhibitors competing for binding at more than one substrate or cosubstrate binding site, sometimes termed bisubstrate inhibitors, can be useful. Examples can be found in Wang et al, BIOCHEMISTRY-US: 15676-15683 2001; and Lerner et al, ANGEW CHEM INT EDIT 40, 4040-4041, 2001.

In the case of FIH and other 2OG oxygenases bisubstrate inhibitors may be useful since features of 2OG binding may be present in more than one enzyme whereas the CAD substrate is unique. Thus, inhibitors that bind to both binding sites may show improved selectivity over those that bind to the 2OG binding site only. The structural analyses enable the identification of such bisubstrate inhibitors. The 2OG and CAD binding sites are linked to each other via a 'channel' extending from the 2-oxo group of 2OG (or NOG) to the beta-carbon Asn-803 in the FIH.Fe.2OG/NOG.HIF(CAD) complexes. In the structures this 'channel' either appears empty but may be occupied by water molecules. The distance from the C of the 2-oxo group of 2OG to the beta-C of Asn-803 is ca. 6 Å. The distance from the 3-C of 2OG to the beta-C of Asn-803 is ca. 6.6 Å. The information from the structural analyses enables the identification of bisubstrate inhibitors, including the following:

These are compounds of formulae (II) to (IV) as defined above except that they are modified such that they can also bind into the 2OG binding pocket as defined by the crystallographic information. Thus, either R^2 or R^1 is modified such that they can bind into the 2OG binding pocket. The modification takes the form such that the general formula of R^1 or R^2 is A-X where X is as defined above and A links X to (II). A is of appropriate length such that X can bind to formula 1 the residues of the 5-

carboxylate of 2OG as discussed above under the heading Inhibitors Exploiting the 2OG binding sites.

More generally bi-substrate inhibitors of FIH can have the formula:



where X is as defined above, B is a linker group and C is an entity binding to part of the CAD binding site of FIH, in general CONH_2 .

B is typically a polymethylene group, generally having 6 to 8 carbon atoms or
10 an equivalent group where one or more of the carbon atoms is replaced by a heteroatom, notably O, S or N and can be functionalised, for example with thiol, alcohol, carboxylate, hydroxamic acid or oxalate to mediate Fe binding. It is preferably 6 to 8 carbon atoms long or its equivalent. Alternatively, B is a linking group which possesses a ring, preferably of 5 to 7 members to which C is attached.

15

Inhibitors that bind to the 2OG binding site or part thereof and the peptide substrate

Another class of inhibitors bind to the enzyme-substrate complex, i.e. to FIH.Fe(II).HIF(CAD). The structural analyses enable the identification of such inhibitors. As described above 2OG and CAD binding sites are linked to each other
20 via a 'channel' extending from the 2-oxo group of 2OG (or NOG) to the beta-carbon of Asn-803 in the FIH.Fe.2OG/NOG.HIF(CAD) complexes.

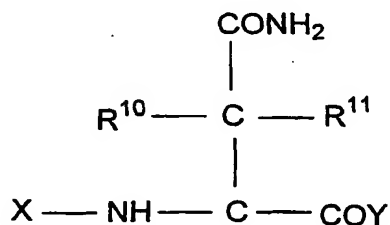
Inhibitors of this type may be defined as $X-[B]-[E]$ where X is as defined above, B is a linker group such as defined above and E is an entity binding to part of the CAD when bound to HIF. E may bind to the backbone carbonyl oxygen of Asn-
25 803 of CAD and to the NH_2 group of the primary amide of Asn-803.

Mechanism based inhibitors

Another class of inhibitors is based upon substrate analogues that can undergo part of the catalytic cycle but either stall at an intermediate stage or cause an aberrant
30 reaction resulting in damage or inhibition. The observation that FIH catalyses hydroxylation of Asn-803 at the beta-position together with the structural analyses enables the design of such inhibitors. Such compounds include analogues of the

substrates (inhibitors) in which Asn-803 is replaced with an analogue which does not undergo oxidation such as beta-fluoro- asparagine, beta-di-fluoro- asparagine, beta-methyl- asparagine, beta-dimethyl- asparagine derivatives. Alternatively derivatives that undergo oxidation to give an agent that can be oxidised to give an inactivating group such as an epoxide or metal chelating group may prepared (such mechanism based inhibitors are sometimes referred to as suicide inhibitors). In the case of FIH they include alpha-beta-dehydroasparagine and beta-methylene asparagine.

These include a compound having the formula



10

wherein X represents a valine residue or an analogue thereof and Y represents an alanine residue or an analogue thereof, R^{10} is fluorine or $\text{C}_1 - \text{C}_3$ alkyl, especially methyl, and R^{11} is fluorine, $\text{C}_1 - \text{C}_3$ alkyl or hydrogen i.e. the specified residue is β -mono- or di-fluoroasparagine or β -mono- or di-methylasparagine.

15

Alternatively, the compound above may be desaturated, i.e. is an alpha/beta dehydroamino acid (R^{11} not present) or R^{10} and R^{11} may be replaced by a methylene group, i.e. the residue is α , β -dehydro-asparagine or β -methylene asparagine.

20

If desired the valine residue is connected to one or more units of the peptide DESGLPQLTSYDCE - in the order given e.g. to glutamic acid (E) alone or to, for aspartic acid (D) - cysteine (C) - glutamic acid (E)-, or a longer chain such as PQLTSYDCE -.

25

For the compounds of this invention suitable aryl rings include phenyl and naphthalenyl, which may be further functionalised or fused to other ring systems. Suitable heterocyclic rings include thiophene, pyridine, quinoline, isoquinoline, pyrimidine, pyrazine, pyrone, chromone, coumarin, indole, isoindole, indolizine, benzofuran, pyridazine, purine, oxazole, pyrazole, isothiazole, pyrrolidine,

piperidine, indoline, benzothiaphen, morpholine, benzimidazole, azepine, azacine, azoine, oxepine, oxocine, oxoine, piperazine, oxazine, thiazine, thiepine, thiocine, thioine, furan, imidazole, azole, diazole, triazole and tetrazole ring systems that may be functionalised or fused to other ring systems.

5 The said alkyl and aryl groups and chains are typically functionalised by alcohol, fluorine, thiol, a carboxylic acid, phosphonic or phosphinic acid, sulphonic acid or other chelating group, in the case of the chains typically via an alkyl group. In the formulae described herein, a branched or straight C₁ to C₆ alkyl chain may be a methyl, ethyl, propyl, butyl, iso-butyl, *tert*-butyl, pentyl, neopentyl, *tert*-pentyl or a
10 primary, secondary or tertiary hexyl group. Preferably the alkyl groups are methyl, the preferred heterocyclic rings are pyrrolidine and tetrahydropyran and the preferred aromatic rings are benzene, naphthalene and pyridine.

 The compounds which are acids can be present in the form of salts, such as sodium salts.

15 The crystal structure of FIH also allows identification of those residues involved in asparaginyl hydroxylase activity of FIH. The crystal structures may therefore be used to design modified FIH, for example, which has reduced or no asparaginyl hydroxylase activity, for example, by mutation of critical residue within the active site. In the alternative, those residues involved in substrate binding can be
20 identified and modified, for example, to allow the asparaginyl hydroxylase to accept other substrates than HIF. For example, by enlarging or decreasing the asparagine binding pocket. Such modified asparaginyl hydroxylases can then be produced using standard techniques. The expected activity can then be assayed as described in more detail below, for example, to identify whether the hydroxylase activity with respect
25 to HIF has been reduced or removed, or alternatively, to assess the asparaginyl activity or binding in respect to other substrates.

 Compounds which have been identified in accordance with the present invention can be further analysed in assays to monitor for activity of the asparagine hydroxylase enzyme directly. Agents which inhibit or reduce HIF asparagine
30 hydroxylase activity reduce hydroxylation of HIF- α and lead to an increase in the interaction with P300 and in particular the CH1 domain and thus transcriptional activation. This in turn will lead to the activation of systemic local defences against

hypoxia or ischaemia that may include the promotion of angiogenesis, erythropoiesis, energy metabolism, inflammation, vasomotor function and will also affect apoptotic/proliferative responses.

We describe below in more detail a number of different assays that may be carried out to assay the activity of modulators of HIF hydroxylase activity or of FIH identified in accordance with the present invention and in particular of asparagine hydroxylase activity, or which affect regulation of HIF- α interaction with p300 in a cell and hence which affect HIF mediated activity. Some of these assays utilise HIF polypeptides, and HIF asparagine hydroxylases. Typically, the assays may utilise a human HIF asparagine hydroxylase such as FIH or a fragment or variant of a human HIF asparagine hydroxylase. These components are described in more detail below. Each of these components, where required, may be provided either in purified or unpurified form, for example, as cellular extracts or by purification of the relevant component from such extracts. Alternatively, the relevant component can be expressed using recombinant expression techniques and purified for use in the assay. Alternatively, the components may be expressed recombinantly in a cell for use in cell based assays.

Typically, a polynucleotide encoding the relevant component is provided within an expression vector. Such expression vectors are routinely constructed in the art and may for example involve the use of plasmid DNA and appropriate initiators, promoters, enhancers and other elements, such as for example polyadenylation signals which may be necessary and which are positioned in the correct orientation in order to allow full protein expression. Suitable vectors would be very readily apparent to those of skill in the art, such as those described in more detail herein with reference to the HIF hydroxylases. Promoter sequences may be inducible or constitutive promoters depending on the selected assay format. The promoter may be tissue specific. Examples of promoters and other flanking sequences for use in the expression vectors are described in more detail herein with reference to the HIF hydroxylases of the invention and in particular to the human HIF hydroxylases.

HIF Polypeptides and Peptide Analogues

The assays of the present invention may use a substrate of a HIF asparagine hydroxylase and in particular an asparagine containing substrate of the enzyme. In particular, such substrates may be used in assays to monitor for the activity of a modulator of HIF asparagine hydroxylase activity. The substrate may be a HIF polypeptide or peptide analogue thereof. Typically, a HIF polypeptide will be used as the substrate.

Any suitable substrate in which an asparagine residue is hydroxylated by a FIH may be used. In preferred embodiments of the invention, such a substrate is a HIF polypeptide such as a HIF-1 α or HIF-2 α subunit protein or fragment of either or peptide analogue of the subunit or fragment. Preferably, the HIF- α peptide conveys an oxygen regulated response. Preferably, the HIF- α peptide has a CAD domain and is capable of oxygen regulated interaction with p300 and downstream transcriptional activation. Preferably, such HIF- α peptides are capable of interacting with the p300 CH1 domain. Preferably, such HIF polypeptides, fragments or peptide analogues incorporate an asparagine residue equivalent to Asn 803 defined with reference to HIF-1 α . The asparagine equivalent to Asn 803 of HIF-1 α may be determined by aligning the HIF variant, fragment or analogue to the sequence of HIF-1 α to obtain the best sequence alignment and identifying thereby the asparagine equivalent to Asn 803 of HIF-1 α .

A HIF polypeptide may be of eukaryotic origin, in particular a human or other mammalian, HIF- α subunit protein or fragment thereof. Alternatively, the polypeptide may be of *C. elegans* origin. In those assays which monitor for hydroxylation of HIF- α through its interaction with p300, the HIF polypeptide has the ability to bind to a wild type full length p300 protein or a fragment thereof comprising the CH1 domain. Preferably, such binding is able, in a hypoxic cellular environment, to activate transcription.

A number of HIF α subunit proteins have been cloned. These include HIF-1 α , the sequence of which is available as Genbank accession number U22431, HIF-2 α , available as Genbank accession number U81984 and HIF-3 α , available as Genbank accession numbers AC007193 and AC079154. These are all human HIF α subunit proteins and all may be used in the invention. HIF- α subunit proteins from

other species, including murine HIF-1 α (accession numbers AF003695, U59496 and X95580), rat HIF-1 α (accession number Y09507), murine HIF-2 α (accession numbers U81983 and D89787) and murine HIF-3 α (accession number AF060194) may also be used in the invention.

5 One HIF- α protein of particular interest is the *C.elegans* HIF- α subunit protein. The *C.elegans* system may be used in assays of the present invention.

There are a number of common structural features found in the two HIF- α subunit proteins identified to date. Some of these features are identified in O'Rourke *et al* (1999, J. Biol. Chem., 274; 2060-2071) and may be involved in the trans-
10 activation functions of the HIF- α subunit proteins. One or more of these common structural features are preferred features of the HIF polypeptides.

Variants of the above HIF- α subunits may be used, such as synthetic variants which have at least 45% amino acid identity to a naturally occurring HIF- α subunit (particularly to a human HIF- α subunit such as, for example HIF-1 α), preferably at
15 least 50%, 60%, 70%, 80%, 90%, 95% or 98% identity. Such variants may include substitutions or modifications as described above with respect to HIF hydroxylases. Amino acid activity may also be calculated as described above with reference to HIF hydroxylases.

HIF fragments may also include non-peptidyl functionalities and may be
20 optimised for assay purposes such that the level of identity is lowered. Such functionalities may be covalently bound such as sugars or non-covalently bound such as metal ions.

HIF α polypeptides as described herein may be fragments of the HIF- α subunit protein or variants as described above, provided that said fragments retain the
25 ability to interact with a wild-type p300 CH1 domain. When using proteinogenic amino acid residues, such fragments are desirably at least 20, preferably at least 40, 50, 75, 100, 200, 250 or 400 amino acids in size. Desirably, such fragments include asparagine 803.

Cell based assays of the present invention may involve upregulation of an
30 endogenous HIF- α or expression of a HIF- α by recombinant techniques and in particular of HIF-1 α .

Assay Methods

The present invention provides an assay method for an agent identified as a modulator of asparagine hydroxylation of hypoxia inducible factor. The method comprises contacting a HIF asparagine hydroxylase and a test substance in the presence of a substrate of the hydroxylase under conditions in which asparagine hydroxylation occurs in the absence of the test substrate and determining asparagine hydroxylation of the substrate. In an alternative assay, HIF asparagine hydroxylase and the test substance are contacted in the presence of the substrate of the hydroxylase under conditions in which hydroxylation does not occur in the absence of the test substrate. Determination of any asparagine hydroxylation is monitored to identify whether the agent actively acts as a promoter of asparagine hydroxylase.

FIH has been found to hydroxylate HIF- α at an asparagine residue within the CAD domain. This hydroxylation mediates p300 binding and in particular, reduces p300 binding. Such binding leads to transcriptional activation. This interaction and activation may also be used as the basis for an assay of the invention.

Such assays of the present invention may be used to assay the activity of inhibitors of HIF asparagine hydroxylase activity and are thus preferably carried out under conditions under which asparagine hydroxylation would take place in the absence of the test substance. The assays of the invention may also be used to assay the activity of inhibitors which are specific for HIF asparagine hydroxylases and which do not have activity or are less active with other hydroxylases, for example, such as HIF prolyl hydroxylases or other asparagine/aspartamic acid hydroxylases. The assays of the invention may also be used to assay the activity of hydroxylase modulators, such as HIF prolyl hydroxylase inhibitors which are not expected to have activity on FIH based on structural modelling studies, and hence may be used to identify inhibitors which are specific for prolyl hydroxylase.

Methods for monitoring modulation

The precise format of any of the screening or assay methods of the present invention may be varied by those of skill in the art using routine skill and knowledge. The skilled person is well aware of the need to additionally employ appropriate controlled experiments. The assays of the present invention may involve monitoring

for asparagine hydroxylation of a suitable substrate, monitoring for the utilisation of substrates and co-substrates, monitoring for the production of the expected products between the enzyme and its substrate. Assay methods of the present invention may also involve screening for the direct interaction between components in the system.

5 Alternatively, assays may be carried out which monitor for downstream effects such as binding of HIF by p300 and downstream effects mediated by HIF such as HIF mediated transcription using suitable reporter constructs or by monitoring for the upregulation of genes or alterations in the expression patterns of genes known to be regulated directly or indirectly by HIF.

10 Various methods for determining hydroxylation are known in the art and are described and exemplified herein. Any suitable method may be used for determining activity of the HIF hydroxylase such as by substrate or co-substrate utilization, product appearance such as peptide hydroxylation or downstream effects mediated by hydroxylated or non-hydroxylated products.

15 Assays may be carried out to monitor directly for hydroxylation of the relevant asparagine residue or another position. Alternatively, assays may be carried out to monitor for depletion of co-factors and co-substrates. Alternatively, such assays may monitor the downstream effects of hydroxylation of HIF or indeed inhibition of hydroxylation of HIF, for example, by monitoring the interaction
20 between HIF and p300 or HIF mediated transcription. Alternatively, reporter gene constructs driven by HIF regulated promoters may be used. Assays are also provided for the identification of enhancers of the activity of the HIF asparagine hydroxylase. Such enhancers may be used to reduce HIF α activity.

In one embodiment, a suitable substrate of the HIF asparagine hydroxylase is
25 provided. This may be HIF- α or a fragment thereof which includes a CAD domain or which includes a residue equivalent to Asn 803 of HIF-1 α . The substrate may not be initially hydroxylated at the Asn 803 position. This may be achieved by providing synthetic polypeptide substrates, or by producing HIF- α polypeptides in bacterial cells, insect cells or mammalian cells or in *in vitro* transcription and translation
30 systems. Alternatively, assays may be carried out over a selected time course such that the substrate is produced during the course of the assay, initially in un-hydroxylated form.

The substrate, enzyme and potential inhibitor compound may be incubated together under conditions which, in the absence of inhibitor provide for hydroxylation of Asn 803, and the effect of the inhibitor may be determined by determining hydroxylation of the substrate. This may be accomplished by any
5 suitable means. Small polypeptide substrates may be recovered and subject to physical analysis, such as mass spectrometry or chromatography, or to functional analysis, such as the ability to bind to p300 (or displace a reporter molecule from p300). Such methods are known as such in the art and may be practiced using routine skill and knowledge. Determination may be quantitative or qualitative. In
10 both cases, but particularly in the latter, qualitative determination may be carried out in comparison to a suitable control, e.g. a substrate incubated without the potential inhibitor.

Inhibitor compounds which are identified in this manner may be recovered and formulated as pharmaceutical compositions.

15 Assays in accordance with the present invention may involve monitoring for the interaction between p300 and HIF. The interaction between HIF and p300 is mediated by hydroxylation of HIF. Transcription and expression of genes known to be upregulated or down regulated by the presence of HIF could be monitored. In particular, upregulation of HIF regulated genes would demonstrate inhibition of
20 asparagine hydroxylation whereas down regulation would suggest enhancement or promotion of asparagine hydroxylation.

In alternative embodiments, reporter constructs may be provided in which promoters mediated by HIF are provided operably linked to a reporter gene. Any suitable reporter gene could be used, such as for example enzymes which may then
25 be used in colorimetric, fluorometric, fluorescence resonance or spectrometric assays.

HIF asparagine hydroxylase is a 2OG dependent oxygenase. In the assay methods described herein, typically the HIF asparagine hydroxylase and the substrate of the hydroxylase are contacted in the presence of a co-substrate, such as 2-oxoglutarate (2OG). The hydroxylase activity of the HIF hydroxylase may be
30 determined by determining the turnover of the co-substrate. This may be achieved by determining the presence and/or amount of reaction products, such as hydroxylated

substrate or succinic acid. The amount of product may be determined relative to the amount of substrate. Typically, in such embodiments the substrate may be an HIF- α polypeptide and, for example, the product measured may be hydroxylated HIF- α polypeptide.

5 Alternatively, the end-point determination may be based on conversion of HIF α or peptide fragments (including synthetic and recombinant peptides) derived from HIF α into detectable products. Peptides may be modified to facilitate the assays so that they can be rapidly carried out and may be suitable for high throughput screening.

10 For example, reverse phase HPLC (C-18 octadecylsilane column), may be used to separate starting synthetic peptide substrates for HIF hydroxylase from the asparagine hydroxylated products, as the latter have a shorter retention time in the column. Modifications of this assay or alternative assays for HIF hydroxylase activity may employ, for example, mass spectrometric, spectroscopic, and/or
15 fluorescence techniques as are well known in the art (Masimirembwa C. *et al* Combinatorial Chemistry & High Throughput Screening (2001) 4 (3) 245-263, Owicki J. (2000) J. Biomol. Screen. 5 (5) 297-305, Gershkovich A *et al* (1996) J. Biochem. & Biophys. Meths. 33 (3) 135-162, Kraaft G. *et al* (1994) Meths. Enzymol. 241 70-86). Fluorescent techniques may employ versions of the substrate
20 modified in such as way as to carry out or optimise spectroscopic or fluorescence assays.

For example, HIF α polypeptide may be immobilised e.g. on a bead or plate, and hydroxylation of the appropriate residue detected using an antibody or other binding molecule which binds the CAD binding domain of HIF α with a different
25 affinity when an asparagine 803 is hydroxylated from when the residue is not hydroxylated. Such antibodies may be obtained by means of standard techniques which are well known in the art, e.g. using a hydroxylated HIF α peptide.

Binding of a molecule which discriminates between the hydroxylated and non-hydroxylated form of a HIF α polypeptide may be assessed using any technique
30 available to those skilled in the art, which may involve determination of the presence of a suitable label.

Assay methods of the present invention may also take the form of an *in vivo* assay. The *in vivo* assay may be performed in a cell line such as a yeast strain in which the relevant polypeptides or peptides are expressed from one or more vectors introduced into the cell.

5

In vivo assays

The assays may be carried out using cell based, organ based or whole animal assays conducted *in vivo*. Such assays may utilize the endogenous expression of the HIF hydroxylase nucleotides and/or polypeptides. In other forms of the invention, upregulation of specific endogenous HIF hydroxylases may be achieved by
10 stimulators of the expression thereof. Such stimulators may be growth factors or chemicals that upregulate specific HIF asparagine hydroxylases. In another form of the assay, nucleotide constructs may be introduced into cells or transgenic animals to increase production of one or more specific HIF asparagine hydroxylases.

15 HIF complexed with p300 activate hypoxia response elements that are found in the promoters and/or enhancers of endogenous genes that are regulated by the said HIF complexes. Such hypoxia response elements may also be isolated and operationally linked to reporter genes so as to assay the activity of the HIF complex through detection and/or quantitation of the reporter gene or its product. Therefore in
20 a further form of the invention the activity of a HIF- α polypeptide that is regulated by HIF asparagine hydroxylase will be assayed by measuring the effects of the HIF complex on the expression of an endogenous gene or reporter gene that is functionally linked to a HIF binding hypoxia response element. Examples of endogenous genes that are regulated in this way are to be found in the role of the aryl
25 hydrocarbon nuclear translocator (ARNT) in hypoxic induction of gene expression, see for example, Studies in ARNT-deficient cells. S.M. Wood, J.M. Gleadle, C.W. Pugh, O. Hankinson, P.J. Ratcliffe. Journal of Biological Chemistry 271 (1996) 15117-15123, and Hypoxia inducible expression of tumor-associated carbonic anhydrases, C.C. Wykoff, N.J.P. Beasley, K.J. Turner, J. Pastorek, A. Sibtain. G.D.
30 Wilson, H. Turley, K. Talks, P.H. Maxwell, C.W. Pugh, P.J. Ratcliffe, A.L. Harris. Cancer Research 60 (2000) 7075-7083. Examples include but are not limited to glucose transporter isoform 1, phosphoglycerate kinase-1, carbon anhydrase isoform

9, vascular endothelial growth factor. Each of said genes contains one or hypoxia response elements that may be isolated and operationally linked as single or multiple copies to a reporter gene for the measurement of activity of a HIF- α polypeptide that varies in accordance with the activity of a HIF hydroxylase.

5 The activity of genes or gene products that are regulated by a HIF- α polypeptide in accordance with the activity of a HIF hydroxylase affects cellular, organ, and animal physiology. Assays that utilise a specific functional response that is regulated in accordance with the activity of a HIF- α polypeptide in accordance with the activity of a HIF hydroxylase may be used. Such responses include the
10 uptake rate of glucose or glucose analogues that are not metabolized, the growth of blood vessels by angiogenesis, the activity of a carbonic anhydrase enzyme. It is recognised that many other responses that operate at a cellular or systemic level are controlled by the activity of a HIF- α polypeptide in accordance with the activity of a HIF hydroxylase and may be utilized as assays of the said HIF hydroxylase activity
15 in further aspects of the invention.

A HIF- α polypeptide that is a substrate for a HIF hydroxylase may be fused to a further polypeptide so as to cause the activity of the said HIF hydroxylase to regulate the activity of the fusion peptide. Accordingly a further form of the invention provides for the assay of the activity of a fusion polypeptide. In the
20 preferred form such a fusion polypeptide may contain the whole or part of a HIF- α polypeptide, particularly including Asn 803, or the CAD domain. The Gal4 DNA binding domain including the amino acids 1-143 together with the Gal binding upstream activating sequence (UAS) is an example of such a transcription factor and cognate DNA response element whose operation can be assayed by those skilled in
25 the art.

Selectivity

It may also be advantageous to modulate HIF asparagine hydroxylase selectively, as a single target, or in selected hydroxylase groups as well as an entire
30 family. Agents which modulate HIF asparagine hydroxylase activity are therefore preferably specific i.e. they have an increased or enhanced effect on a HIF asparagine hydroxylase relative to other 2OG dependent oxygenases.

Assay methods as described herein may therefore further comprise contacting the test compound with one or more 2OG dependent oxygenases under conditions in which said 2OG dependent oxygenases are normally active and determining activity of said oxygenases. A difference in activity in the presence relative to the absence of test compound is indicative of the test compound modulating the activity of the one or more 2OG dependent oxygenases.

A test compound which provides increased or enhanced modulation of a HIF asparagine hydroxylase, relative to the one or more 2OG dependent oxygenases shows selectivity or specificity for the HIF hydroxylase.

2OG dependent oxygenases may include for example, clavaminte synthase, Alk B deacetoxycephalosporin C synthase, collagen-prolyl-4-hydroxylase, collagen prolyl-3-hydroxylase, lysyl hydroxylase, aspartyl hydroxylase, phytanoyl coenzyme A hydroxylase or gamma-butyrobetaine hydroxylase. 2OG dependent oxygenases may be mammalian, preferably human polypeptides.

The invention provides for the use of such selective inhibitors of HIF asparagine hydroxylases in the manufacture of a medicament for the treatment of a condition associated with reduced HIF activity.

Therapeutic Applications

A compound, substance or agent which is found to have the ability to affect the hydroxylase activity of a HIF asparagine hydroxylase, or the compounds referred to herein as FIH inhibitors has therapeutic and other potential in a number of contexts. For therapeutic treatment, such a compound may be used in combination with any other active substance, e.g. for anti-tumour therapy another anti-tumour compound or therapy, such as radiotherapy or chemotherapy.

An agent identified using one or more primary screens (e.g. in a cell-free system) as having ability to modulate the HIF α asparagine hydroxylation activity of a HIF hydroxylase may be assessed further using one or more secondary screens. A secondary screen may involve testing for an increase or decrease in the amount of HIF- α or HIF activity, for example as manifest by the level of a HIF target gene or process present in a cell in the presence of the agent relative to the absence of the agent.

A HIF hydroxylase or a HIF polypeptide may be used in therapies which include treatment with full length polypeptides or fragments thereof, or otherwise modified polypeptides (e.g. to enhance stability or ensure targeting, including in conjunction with other active agents such as antibodies. For example, mutation of
5 HIF-1 α to replace Asn 803 with another amino acid residue may prevent hydroxylation and thus promote interaction of HIF- α with p300 and stimulate transcriptional activation.

Generally, an agent, compound or substance which is a modulator according to the present invention is provided in an isolated and/or purified form, i.e.
10 substantially pure. This may include being in a composition where it represents at least about 90% active ingredient, more preferably at least about 95%, more preferably at least about 98%. Any such composition may, however, include inert carrier materials or other pharmaceutically and physiologically acceptable excipients, such as those required for correct delivery, release and/or stabilisation of the active
15 agent. Typically, the concentration in such compositions is 0.1 to 50%, generally 0.5 to 20%, especially 1 to 10% by weight based on the weight of the composition. As noted below, a composition according to the present invention may include in addition to an modulator compound as disclosed, one or more other molecules of therapeutic use, such as an anti-tumour agent.

20

Products obtained by assays of the invention

The invention further provides compounds obtained or identified by methods of the present invention, and compositions comprising said compounds, such as pharmaceutical compositions wherein the compound is in a mixture with a
25 pharmaceutically acceptable carrier or diluent. The carrier may be liquid, e.g. saline, ethanol, glycerol and mixtures thereof, or solid, e.g. in the form of a tablet, or in a semi-solid form such as a gel formulated as a depot formulation or in a transdermally administerable vehicle, such as a transdermal patch.

The invention further provides a method of treatment which includes
30 administering to a patient an agent which interferes with the hydroxylation of the asparagine target residue of an HIF α polypeptide by a HIF hydroxylase. Such agents may include inhibitors of asparagine hydroxylase activity. The invention also

provides a method of treatment which includes administering to a patient a compound as defined above.

The therapeutic/prophylactic purpose may be related to the treatment of a condition associated with reduced or suboptimal or increased HIF levels or activity, or conditions in which have normal HIF levels, but where an modulation in HIF activity such as an increase or decrease in HIF activity is desirable such as:

- (i) ischaemic conditions, for example organ ischaemia, including coronary, cerebrovascular and peripheral vascular insufficiency. The therapy may be applied in two ways; following declared tissue damage, e.g. myocardial infarction (in order to limit tissue damage), or prophylactically to prevent ischaemia, e.g. promotion of coronary collaterals in the treatment of angina.
- (ii) wound healing and organ regeneration
- (iii) auto-, allo-, and xeno- transplantation.
- (iv) systemic blood pressure
- (v) cancer; HIF α is commonly up-regulated in tumour cells and has major effects on tumour growth and angiogenesis.
- (vi) inflammatory disorders.
- (vii) pulmonary arterial blood pressure, neurodegenerative disease.

Modulating HIF prolyl hydroxylase activity in a person, an organ, or a group of cells may be exploited in different ways to obtain a therapeutic benefit:

- (a) Non cell autonomous: The HIF system is used by cells to influence the production of substances which signal to other cells. These signals may then have effects at (i) a distant site (for example erythropoietin acts on the bone marrow) or (ii) locally (angiogenic growth factors increase the local formation of blood vessels). Manipulating non cell autonomous behaviour via altering hydroxylase activity is therefore useful in the treatment of anaemia, and local ischaemia, for example in the eye, brain, heart and limbs. Many other signals that are involved in aspects of physiological homeostasis may be, or are known to be, adjusted by HIF activation. Consequently altering HIF prolyl hydroxylase activity may be used to potentiate or initiate a helpful response for a therapeutic benefit, or to prevent or ameliorate a harmful response. For example, this approach can be used to alter appetite, or blood pressure in the systemic or pulmonary beds.

(b) Cell autonomous: the HIF system is also used by cells to regulate cellular metabolism, and decisions concerning differentiation, proliferation and apoptosis. Therefore manipulating the HIF system can be used to alter the viability and behaviour of cells. An increase in cell viability can be achieved by increasing HIF activation, for example in an ischaemic tissue. This approach can also be used in improving pancreatic beta cell viability as a way of ameliorating diabetes, or of improving the viability or function of a group or groups of neurons in Parkinson's disease, motoneurone disease or forms of dementia. In a different approach, the HIF signal can be manipulated to prevent a group of cells proliferating, or to promote its death or differentiation. For example transient activation of the HIF system in a malignant tumour can be used to provoke death of a substantial number of tumour cells.

Pharmaceutical Compositions

In various further aspects, the present invention thus provides a pharmaceutical composition, medicament, drug or other composition for such a purpose, the composition comprising one or more agents, compounds or substances as described herein, including HIF asparagine hydroxylase inhibitors, or one or more compounds of formula (A) to (F) or derivatives thereof, the use of such a composition in a method of medical treatment, a method comprising administration of such a composition to a patient, e.g. for treatment (which may include preventative treatment) of a medical condition as described above, use of such an agent compound or substance in the manufacture of a composition, medicament or drug for administration for any such purpose, e.g. for treatment of a condition as described herein, and a method of making a pharmaceutical composition comprising admixing such an agent, compound or substance with a pharmaceutically acceptable excipient, vehicle or carrier, and optionally other ingredients.

In one embodiment the method for providing a pharmaceutical composition may typically comprise:

- (a) identifying an agent in accordance with the invention; and
- (b) formulating the agent thus identified with a pharmaceutically acceptable excipient.

The agent may be used as sole active agent or in combination with one another or with any other active substance, e.g. for anti-tumour therapy another anti-tumour compound or therapy, such as radiotherapy or chemotherapy.

Whatever the agent used in a method of medical treatment of the present invention, administration is preferably in a "prophylactically effective amount" or a "therapeutically effective amount" (as the case may be, although prophylaxis may be considered therapy), this being sufficient to show benefit to the individual. The actual amount administered, and rate and time-course of administration, will depend on the nature and severity of what is being treated. Prescription of treatment, e.g. decisions on dosage etc, is within the responsibility of general practitioners and other medical doctors.

An agent or composition may be administered alone or in combination with other treatments, either simultaneously or sequentially dependent upon the condition to be treated, e.g. as described above.

Pharmaceutical compositions according to the present invention, and for use in accordance with the present invention, may include, in addition to active ingredient, a pharmaceutically acceptable excipient, carrier, buffer, stabiliser or other materials well known to those skilled in the art. In particular they may include a pharmaceutically acceptable excipient. Such materials should be non-toxic and should not interfere with the efficacy of the active ingredient. The precise nature of the carrier or other material will depend on the route of administration, which may be oral, or by injection, e.g. cutaneous, subcutaneous or intravenous. The compositions will typically be sterile.

Pharmaceutical compositions for oral administration may be in tablet, capsule, powder or liquid form. A tablet may include a solid carrier such as gelatin or an adjuvant. Liquid pharmaceutical compositions generally include a liquid carrier such as water, petroleum, animal or vegetable oils, mineral oil or synthetic oil. Physiological saline solution, dextrose or other saccharide solution or glycols such as ethylene glycol, propylene glycol or polyethylene glycol may be included.

For intravenous, cutaneous or subcutaneous injection, or injection at the site of affliction, the active ingredient will be in the form of a parenterally acceptable aqueous solution which is pyrogen-free and has suitable pH, isotonicity and stability.

Those of relevant skill in the art are well able to prepare suitable solutions using, for example, isotonic vehicles such as Sodium Chloride Injection, Ringer's Injection, Lactated Ringer's Injection. Preservatives, stabilisers, buffers, antioxidants and/or other additives may be included, as required.

5 Liposomes, particularly cationic liposomes, may be used in carrier formulations. Examples of techniques and protocols mentioned above can be found in Remington's Pharmaceutical Sciences, 16th edition, Osol, A. (ed), 1980.

 The substance or composition may be administered in a localised manner to a particular site or may be delivered in a manner in which it targets particular cells or
10 tissues, for example using intra-arterial stent based delivery.

 Targeting therapies may be used to deliver the active substance more specifically to certain types of cell, by the use of targeting systems such as antibody or cell specific ligands. Targeting may be desirable for a variety of reasons, for example if the agent is unacceptably toxic, or if it would otherwise require too high a
15 dosage, or if it would not otherwise be able to enter the target cells.

 In a further embodiment the invention provides for the use of an agent of the invention in the manufacture of a medicament for the treatment of a condition associated with increased or decreased HIF levels or activity. The condition may, for example, be selected from the group consisting of ischaemia, wound healing, auto-
20 allo-, and xeno- transplantation, systemic high blood pressure, cancer, and inflammatory disorders.

Examples

25 **Example 1**

 The position on Asn803 of human HIF-1 α that is hydroxylated was identified as described in the following. cDNA sequences encoding FIH-1 were cloned into the pET28a(+) vector (from Novagen) to yield FIH-1 protein with an N-terminal His₆tag to facilitate purification. Purification of crude material by nickel affinity
30 chromatography, followed by thrombin cleavage of the His₆ tag, and size exclusion chromatography (Superdex S75) yielded >95% pure protein by SDS-PAGE analysis. Mass spectrometry confirmed the identity of the isolated species. The 19-residue

peptide comprising amino acids 788-806 of human HIF-1 α was modified by aerobic incubation with FIH-1 FIH (Hewitson et al., J BIOL CHEM 277 (29): 26351-26355, 2002) in the presence of ascorbate, DTT, catalase, 2-oxoglutarate, and iron(II) for 30 minutes at 37°C. The reaction was quenched by cooling to 4°C and addition of an equal volume of methanol. Precipitate was removed by centrifugation and the supernatant purified by HPLC using a Jupiter C4 column (15cm x 4.6mm). Peptide was eluted using a gradient of acetonitrile in 0.1% trifluoroacetic acid, freeze-dried from the HPLC solvent for amino acid and mass spectrometric analyses. The sample was freeze-dried a second time from D₂O in preparation for NMR analysis.

Catalytic FIH-1 mediated hydroxylation of a synthetic 19 residue peptide corresponding to residues 788-806 of HIF-1 α was confirmed by mass spectrometric analysis of HPLC purified material: Native peptide 19mer $[M+2H]^{2+} = 1026.67\text{Da}$, modified peptide 19mer $[M+2H]^{2+} = 1034.61\text{Da}$, a mass difference of +8Da of the doubly charged ions, corresponding to +16Da in the peptide (oxygen). N-Terminal Edman degradation of the product peptide gave the following sequence: DESGLPQLTSYDCEVxA, where x was not asparagine. The peak from this (16th) cycle of Edman degradation ran to a similar position as the β -hydroxyasparagine standard. Acid hydrolysis of the modified peptide followed by amino acid analysis showed the presence of β -hydroxyaspartic acid only.

Both ^1H and ^{13}C chemical shift changes between the 19mer peptide substrate and the HPLC purified incubation product were assessed by 2D ^1H - ^{13}C HSQC experiments. In the substrate a grouping of four β -CH₂ resonances were assigned as belonging to Asp-1, Tyr-11, Asp-12 and Asn-16 according to their ^1H and ^{13}C shifts (Evans, J. N. S. (1995) Biomolecular NMR Spectroscopy, Oxford University Press, Oxford, UK). In the product it was clear from both the 2D HSQC and the 1D proton spectra that only three of these four resonances are present. Comparison of the two spectra indicates that the signal assigned to the Asn-16 β -carbon (at δH 2.813 and 2.695ppm and δC 37.40ppm in the substrate) has disappeared, consistent with hydroxylation of the asparagine residue at its β -carbon. The resonances due to the two aspartic acid residues had shifted slightly, presumably due to changes in the protonation state, and now occur at a similar ^1H chemical shift as the β -protons of the

asparagine in the substrate. A difference in the oxidation state of the cysteine between the two samples is unlikely given the near identical chemical shifts for the cysteinyl β -carbon and hydrogens. The change from a double doublet to a single doublet for the β -hydrogen of the hydroxylated residue also rules out any possibility the observed alterations in the NMR spectrum are due to aggregation. Two new resonances have appeared in the product spectrum at δ H 4.913 ppm and δ C 56.26 ppm and at δ H 4.654 ppm and δ C 72.22 ppm. These resonances correlate with one another in the 2D COSY spectrum and share a ^1H - ^1H coupling constant of 2.4 Hz and are therefore assigned as the CH^α - CH^β of the hydroxylated asparagine. The appearance of these resonances also coincides with the disappearance of the δ H 4.706 ppm and δ C 51.43 ppm resonances observed in the substrate spectra, which is therefore assigned as the CH^α of the parent asparagine prior to modification. Comparison of the CH^α - CH^β coupling constant of 2.4 Hz observed for the hydroxylated Asn-803, with literature values implied the *threo* isomer is produced.

In summary of the above ^1H -NMR experiments: The HSQC experiments gave direct evidence for hydroxylation occurring at the β -carbon of the target asparagine, with the hydroxylated β -carbon appearing significantly deshielded (at 72.22 ppm) and the adjacent α -carbon deshielded to a lesser extent (at 56.26 ppm) relative to the parent asparagine. Changes of these magnitudes in the ^{13}C chemical shifts are inconsistent with hydroxylation of the side-chain nitrogen, but consistent with hydroxylation at the β -carbon. Further, the ^{13}C spectrum of free DL-*threo*- β -hydroxyasparagine (this study), has resonances at 58.63 ppm and 73.85 ppm corresponding to α - and β -carbons. The product assignment is also consistent with ^1H -NMR chemical shifts of the α - and β -hydrogens in the β -hydroxyaspartyl residues in EGF-like domains which are 4.48 ppm and 4.36 ppm respectively (with respect to water at 4.75 ppm) when calcium is absent (Selander et al, Biochemistry 29, 8111-8118). The analysis of the coupling constant reported here suggests that the *threo*-isomer is the one formed on hydroxylation of Asn-803 by FIH-1.

Two reports (Dames et al., (2002) *Proc. Natl. Acad. Sci. U. S. A.* 99, 52715276; Freedman et al, (2002) *Proc. Natl. Acad. Sci. U. S. A.* 99, 53675372) reveal how β -hydroxylation of Asn-803 of HIF-1 α would be damaging to complex formation with

p300. Although the position of hydroxylation was not identified in either report, both imply that hydroxylation at the *pro-S* position of the β -carbon, *i.e.* to give the *threo* (2*S*, 3*S*)-isomer, would interfere with the hydrogen bonding that maintains the α -helical conformation adopted by this part of HIF-1 α , and also create a need for the energetically unfavourable desolvation of the hydroxyl group. A steric clash between the inserted *pro-S* hydroxyl group and Ile-353 (numbering from Dames *et al* (2002) *Proc. Natl. Acad. Sci. U. S. A.* **99**, 52715276) of p300 would disrupt the interaction of the two proteins. Presumably the same mechanism is also used to abrogate the interaction of HIF-2 α and p300. The discovery that it is the beta-position of Asn-803 that is modified and the associated mechanistic implications may be used in the design of compounds that bind to p300 thereby displacing HIF- α and/in the design of inhibitors of FIH (see below); in both cases to enable pro-angiogenic pharmaceutical agents.

Example 2

To obtain an FIH:CAD complex suitable for X-ray analysis without oxidation of the CAD or the Fe^(II), FIH and various CAD fragments from seven to fifty-two residues were co-crystallised with Fe^(II) and 2OG under anaerobic conditions. Structures were also obtained for FIH complexed with Fe^(II) and *N*-oxaloylglycine (NOG, an FIH inhibitor), (anaerobically) and Zn^(II) and NOG (aerobically). These structures were solved by molecular replacement using a model obtained by multiple anomalous dispersion on selenomethionine-substituted apo-FIH. Crystalline FIH:CAD complexes were obtained with CAD₇₈₆₋₈₂₆, Fe^(II) and NOG or 2OG (structures 1 and 2, Table 1), CAD₇₇₅₋₈₂₆ with Zn^(II) and NOG (structure 3). Crystallisation attempts with CAD₇₈₇₋₈₀₆, CAD₈₅₀₋₈₆₂ (HIF-2 α , equivalent to HIF-1 α CAD₈₀₂₋₈₁₄) and CAD₈₀₀₋₈₀₆ did not result in FIH:CAD complexes; solution analyses indicated that CAD fragments shorter than twenty residues are not efficient *in vitro* substrates.

Table 1 Summary of FIH:CAD-fragment complex structures*

Structure No.	Resolution (Å)	Metal	Co-substrate	Co-crystallisation CAD peptide	Site 1 CAD residues resolved	Site 2 CAD residue resolved	R _{free} (%)	R.m.s.d. from Struct. 1 (Å)	PDB ID.
1	2.15	Fe	NOG	HIF-1α 786-826	795-806	812-823	21.3	-	1H2K
2	2.25	Fe	2OG	HIF-1α 786-826	795-806	813-822	21.7	0.149	1H2L
3	2.50	Zn	NOG	HIF-1α 775-826	795-806	813-822	22.5	0.136	1H2M
4	2.84	Fe	2OG	HIF-2α 850-862	-	-	25.7	0.226	1H2N

* Crystalline FIH:CAD complexes were also obtained with Fe^(II), HIF-1α 775-786 and 2OG or NOG.

Methods employed in structural work

Protein expression, purification and crystallisation

FIH, CAD₇₇₅₋₈₂₆ and CAD₇₈₆₋₈₂₆ were prepared as described (Hewitson et al., J BIOL CHEM 277 (29): 26351-26355, 2002). Selenomethionine (SeMet) substituted
5 FIH was produced using a metabolic inhibition protocol and LeMaster media supplemented with 50 mg/l L-selenomethionine. SeMet incorporation was >95 % by ESI-MS. Aerobic crystallisation of SeMet FIH (at 11 mg ml⁻¹) was accomplished by hanging-drop vapour diffusion at 17 °C. The mother liquor consisted of 1.2 M ammonium sulphate, 4 % PEG 400 and 0.1 M Hepes pH 7.5. Crystallisation of
10 FIH:Fe:CAD fragment complexes was accomplished under an anaerobic atmosphere of argon in a Belle Technology glove box (0.3-0.4 ppm O₂) using the same mother liquor and a solution containing FIH (at 11 mg ml⁻¹), Fe²⁺ (1 mM), 2OG/NOG (2 mM) and CAD fragment (1 mM). Crystallisation of FIH:Zn:CAD fragment was accomplished aerobically under similar conditions. Peptides were either synthesised
15 by solid phase peptide synthesis or purchased from Biopeptide Co. (San Diego, USA).

Crystallographic data collection and structure refinement

Crystals were cryocooled by plunging into liquid nitrogen and X-ray data
20 were collected at 100 K using a nitrogen stream. Cryoprotection was accomplished by sequential transfer into a solution containing 1.2 M ammonium sulphate, 3 % PEG 400, 0.1 M Hepes pH 7.5 and 10 % followed by 24 % glycerol. A three-wavelength multiple anomalous dispersion (MAD) dataset was collected to 2.9 Å resolution on beamline 14.2 of the Synchrotron Radiation Source, Daresbury, U.K.
25 Data from crystals of FIH:CAD complexes were collected on beamlines 14.2, 9.6 or 9.5 using ADSC Quantum 4 (14.2 and 9.6) or MarCCD detectors (9.5). All data was processed with the program MOSFLM and the CCP4 suite [Collaborative Computational Project Number 4 *Acta Crystallogr.* D50, 760-763 (1994)]. The crystals belonged to space group *P*4₁2₁2. Six selenium positions were located and
30 phases calculated using the program SOLVE (Terwilliger et al. D55, 849-861, 1999). Density modification, which increased the figure of merit from 0.56 to 0.66, was performed using RESOLVE (Terwilliger *Acta Crystallogr.* D56, 965-972 2000).

An initial model was built using the program O (Jones et al, *Acta Crystallogr.* A47, 110-119, 1991).and refined against the SeMet data (remote wavelength) using the program CNS (Brunger *Acta Crystallogr.* D54, 905-921, 1998). . One cycle of simulated annealing followed by grouped *B*-factor refinement brought the R_{free} to 36.2 %. Following further rebuilding and refinement, which brought the R_{free} to 32.3 %, the model was transferred to the 2.15 Å dataset. Rebuilding and refinement using REFMAC5 including addition of Fe, substrate and solvent molecules, and refinement of TLS parameters brought the conventional *R*-factor to 17.8 % and the R_{free} to 21.3 %. The following residues are missing in the current model: 1-15 and 304-306 of FIH, 786-794, 807-811 and 824-826 of the CAD fragment. According to PROCHECK there are no Ramachandran outliers and 90.7 % of residues have most favourable backbone conformations. For the CAD peptide, 77.8% of residues are in the most favourable region with the remaining 22.2 % in additionally allowed regions.

Other structures were solved by molecular replacement using the coordinates from the 2.15 Å data and refinement using REFMAC5. In all structures electron density for the Fe and 2OG/NOG was visible throughout refinement. Significant positive difference electron density was observed between the iron and the CAD Asn803 β-carbon. Since *B*-factor differences between FIH and CAD imply that the CAD is not at 100 % occupancy, this may represent an alternative binding-mode for the 1-carboxylate 2OG in the absence of substrate although it could also be due to a ligating water molecule, again in the absence of substrate.

Overview of FIH structure

The core of FIH comprises a double-stranded beta-helix (DSBH or jellyroll) motif formed from eight β-strands, β8-β11 and β14-β17. Residues 220-259 form an insert between strands 4 and 5 of the DSBH. The bottom face of the DSBH is flanked by an additional four β-strands from the N-terminal region to form an eight-membered antiparallel β-sheet. The N-terminal strand β1 bisects the face of the DSBH opposite to the active site. The β1 strand has a 360° twist located at a PXXP sequence, in between its interactions with β14 and β2. A similarly positioned β-

strand is found in most 2OG oxygenases, though not always from the same region of the protein. The sheet-helix-sheet motif formed by $\beta 1$, $\alpha 1$ and $\beta 2$ is conserved in all enzymes of this class except proline 3-hydroxylase and a similar fold in this region is found in the related $\text{Cu}^{(II)}$ utilising quercetin 2,3-dioxygenase (QD) (Fusetti et al, 5 STRUCTURE 10 (2): 259-268 2002). The topology of FIH unequivocally defines it as an iron-binding member of the cupin structural family which already includes QD and $\text{Mn}^{(II)}$ utilising Type II phosphomannose isomerase (Clissold, P. M., and Ponting, C. P. (2001) *Trends Biochem. Sci.* 26, 79).

10 Related enzymes to FIH

FIH has significant sequence similarity with the JmjC homology region of the jumonji transcription factors (Clissold, P. M., and Ponting, C. P. (2001) *Trends Biochem. Sci.* 26, 79; Hewitson et al., J BIOL CHEM 277 (29): 26351-26355, 2002). These proteins are members of the cupin structural superfamily and have been 15 implicated in cell growth and heart development. The 2OG oxygenase iron binding residues had been identified in some JmjC domains but not assigned as an iron binding motif. Sequence searches in the light of the FIH structure reveal many JmjC proteins with conserved residues that include both this motif and others, including FIH residues Lys214 and Thr196 that are unusually involved in binding the 5- 20 carboxylate of 2OG. The structure thus reveals that FIH is a one of a large family of iron and 2OG dependent oxygenases involved in the regulation of transcription. Since some of the assigned JmjC domains other than FIH are associated with diseases and particular phenotypes their (e.g.) inhibition may be of therapeutic value. (See e.g. Hu et al, ONCOGENE 20 (47): 6946-6954 OCT 18 2001 and Clissold, P. 25 M., and Ponting, C. P. (2001) *Trends Biochem. Sci.* 26, 79 and references therein).

Table 2. Partial sequence alignment of FIH with a selection of JmjC domain containing proteins. FIH secondary structure is indicated above the alignment. Selected 2OG binding residues found in FIH are indicated by dark triangles under the alignment and the two iron binding residues by light triangles. SWALL accession 30 numbers are indicated on the left of the alignment.

	$\alpha 7$	$\beta 8$	$\beta 9$	$\beta 10$	$\beta 11$				
Hs Q969Q7	FIH	FNNWINKQQ	---GKRGWQ	---LTSNLLI	MEGNTVPAHYEQ	---ONFEAQIKGY	---KRCILFPDP		
Dm Q9VU77		---ELAADR	---VSDLDFAQQ (4)	---PPDAVFWL	DERAVTSMHKDPY	---ENVYCVISGH	---KDFVLIPPH		
Dm Q9W0M3		---ALKEDIS	---IPDYCTI (5)	PGAVDIKAWL	PAGTVSPMHYDEK	---HNLLCQVFGS	---KRIILAAPA		
Hs Q9UPP1		---KIVRKLS	---WVENLWEEC (4)	PNVQKYCLMSVRDSY	TDFHIDFGGT	---SWYHYVLKGE	---KIFYLIRPT		
Ce Q9B167		---RFVQEIS	---MVNRLNEDV (20)	PKVEQFCLL	MAGSYTDFHVDFFGS	---SVYHYHLKGE	---KIFYIAAPT		
Ce Q20367		---RFVQDIS	---MAKRLNSDV (11)	PKIEQICAAAMANSY	TDFHVDFFGT	---SVYFHVFKGE (4)	KIFYIAAPT		
Dm Q9VHH9		---EIVRQID	---WVDVWPKQ (16)	PKVQKYCLMSVKNCY	TDFHIDFGGT	---SVYHYHLRGS (1)	KVFWLIRPT		
Sc P40034		---QNDLVDKIF	---SENGHLEKV (11)	PKVTKYILMSVKDAY	TDFHLDFAGT	---SVYNNVISGQ	---KKFLFPPT		
Rn Q9R153	PASS1	---KTDVFQEVW	---WSDFGFP	---RNGQE	---STLWT	SLGAHTPCHLDSYG	---CNLVEQVQGR	---KRWHLFPPE	
Ce Q9GY14		FEDDLFHYAD	---DKKRPDH	---RWFVN	PARSGTAIHIDPLGTS	WNSLLQGH	---KRWVLIPPI		
Dm Q9V6L0		---TILDVWVKDYN	IQID	VNT	---AYLYF	MMKTFEAWHTEDMDLYS	INLHFGAP	---KRWVVPPE	
Hs Q94877		---TVLDVWEEEC	GISIEK	VNT	---PYLYF	MMKTFEAWHTEDMDLYS	INLHFGEP	---RSWVAIPPE	
Ce Q9U297		---TILEDTNVE	---IKQVNT	---VYLYF	MMKTFEAWHTEDMDLYS	INLHFGAP	---KYWFAISSE		
Dm Q9V333		---TILNLVNTDYN	IID	VNT	---AYLYF	MMKTSFAWHTEDMDLYS	INLHFGAP	---KRWVAIPPA	
Hs O75164		---TILDVVEKESG	ITIEG	VNT	---PYLYF	MMKTSFAWHTEDMDLYS	INLHFGEP	---KSWYSVPPE	
Dm Q9VJ97		FASDWLNEQL	---IQQ	KDDY	---RFVYM	PKNSWTSYHADVFGS	FSWSTNIVGL	---KKWLTMPPG	
Sp O13977		FADDWLNAVY	---IDCESDDF	---RFAYL	SHLT	GLT	PDVYASHSF	SVNLCGV	---KCWLFIIDPK

Hs = Homo sapiens
 Dm = Drosophila melanogaster
 Ce = Caenorhabditis elegans
 Sc = Saccharomyces cerevisiae
 Rn = Rattus norvegicus
 Sp = Schizosaccharomyces pombe

FIH = Factor Inhibiting HIF
 PASS1 = Protein associating with small stress protein

TABLE 2

TABLE 3 - Coordinates for structures 1 to 4Structure 1

Below are the coordinates for structure 1 (the 2.15 Å structure of FIH:Fe(II):NOG:CAD):

```

HEADER      TRANSCRIPTION ACTIVATOR/INHIBITOR      12-AUG-02   1H2K
TITLE       FACTOR INHIBITING HIF-1 ALPHA IN COMPLEX WITH HIF-1 ALPHA
TITLE       2 FRAGMENT PEPTIDE
COMPND      MOL_ID: 1;
COMPND      2 MOLECULE: FACTOR INHIBITING HIF1;
COMPND      3 CHAIN: A;
COMPND      4 ENGINEERED: YES;
COMPND      5 MOL_ID: 2;
COMPND      6 MOLECULE: HYPOXIA-INDUCIBLE FACTOR 1 ALPHA;
COMPND      7 SYNONYM: HIF-1 ALPHA, ARNT INTERACTING PROTEIN,
COMPND      8 MEMBER OF PAS PROTEIN 1;
COMPND      9 CHAIN: S;
COMPND     10 FRAGMENT: C-TERMINAL TRANSACTIVATION DOMAIN FRAGMENT
COMPND     11 RESIDUES 786-826
SOURCE      MOL_ID: 1;
SOURCE      2 ORGANISM_SCIENTIFIC: HOMO SAPIENS;
SOURCE      3 ORGANISM_COMMON: HUMAN;
SOURCE      4 EXPRESSION_SYSTEM: ESCHERICHIA COLI;
SOURCE      5 EXPRESSION_SYSTEM_STRAIN: BL21(DE3);
SOURCE      6 EXPRESSION_SYSTEM_PLASMID: PET28A(+);
SOURCE      7 MOL_ID: 2;
SOURCE      8 ORGANISM_SCIENTIFIC: HOMO SAPIENS;
SOURCE      9 ORGANISM_COMMON: HUMAN;
SOURCE     10 EXPRESSION_SYSTEM: ESCHERICHIA COLI;
SOURCE     11 EXPRESSION_SYSTEM_STRAIN: BL21(DE3);
SOURCE     12 EXPRESSION_SYSTEM_PLASMID: PGEX-GP-1
KEYWDS      FIH, HIF, DSBH, OXYGENASE, TRANSCRIPTION, HYPOXIA,
KEYWDS      2 2-OXOGLUTARATE, ASPARAGINYL HYDROXYLASE, PHOSPHORYLATION
EXPDTA      X-RAY DIFFRACTION
AUTHOR      J.M.ELKINS,K.S.HEWITSON,L.A.MCNEILL,I.SCHLEMMINGER,
AUTHOR      2 J.F.SEIBEL,C.J.SCHOFIELD
REVDAT      1   03-SEP-02 1H2K   0
JRNL        AUTH   J.M.ELKINS,K.S.HEWITSON,L.A.MCNEILL,
JRNL        AUTH 2 I.SCHLEMMINGER,J.F.SEIBEL,C.J.SCHOFIELD
JRNL        TITL   FIH:HIF-FRAGMENT COMPLEXES
JRNL        REF    TO BE PUBLISHED
JRNL        REFN
REMARK      2
REMARK      2 RESOLUTION. 2.15 ANGSTROMS.
REMARK      3
REMARK      3 REFINEMENT.
REMARK      3   PROGRAM      : REFMAC 5.0
REMARK      3   AUTHORS      : MURSHUDOV,VAGIN,DODSON
REMARK      3
REMARK      3   REFINEMENT TARGET : MAXIMUM LIKELIHOOD
REMARK      3
REMARK      3 DATA USED IN REFINEMENT.
REMARK      3   RESOLUTION RANGE HIGH (ANGSTROMS) :   2.15
REMARK      3   RESOLUTION RANGE LOW  (ANGSTROMS) :  18.50
REMARK      3   DATA CUTOFF          (SIGMA(F)) :  NONE
REMARK      3   COMPLETENESS FOR RANGE          (%) :  99.28
REMARK      3   NUMBER OF REFLECTIONS              :  28171
REMARK      3

```

```

REMARK 3 FIT TO DATA USED IN REFINEMENT.
REMARK 3 CROSS-VALIDATION METHOD : THROUGHOUT
REMARK 3 FREE R VALUE TEST SET SELECTION : RANDOM
REMARK 3 R VALUE (WORKING + TEST SET) : 0.18026
REMARK 3 R VALUE (WORKING SET) : 0.17761
REMARK 3 FREE R VALUE : 0.21305
REMARK 3 FREE R VALUE TEST SET SIZE (%) : 7.7
REMARK 3 FREE R VALUE TEST SET COUNT : 2340
REMARK 3
REMARK 3 FIT IN THE HIGHEST RESOLUTION BIN.
REMARK 3 TOTAL NUMBER OF BINS USED : 20
REMARK 3 BIN RESOLUTION RANGE HIGH : 2.150
REMARK 3 BIN RESOLUTION RANGE LOW : 2.205
REMARK 3 REFLECTION IN BIN (WORKING SET) : 1906
REMARK 3 BIN R VALUE (WORKING SET) : 0.222
REMARK 3 BIN FREE R VALUE SET COUNT : 152
REMARK 3 BIN FREE R VALUE : 0.257
REMARK 3
REMARK 3 NUMBER OF NON-HYDROGEN ATOMS USED IN REFINEMENT.
REMARK 3 PROTEIN ATOMS : 2875
REMARK 3 NUCLEIC ACID ATOMS : 0
REMARK 3 HETEROGEN ATOMS : 21
REMARK 3 SOLVENT ATOMS : 194
REMARK 3
REMARK 3 B VALUES.
REMARK 3 FROM WILSON PLOT (A**2) : NULL
REMARK 3 MEAN B VALUE (OVERALL, A**2) : 25.725
REMARK 3 OVERALL ANISOTROPIC B VALUE.
REMARK 3 B11 (A**2) : -0.27
REMARK 3 B22 (A**2) : -0.27
REMARK 3 B33 (A**2) : 0.55
REMARK 3 B12 (A**2) : 0.00
REMARK 3 B13 (A**2) : 0.00
REMARK 3 B23 (A**2) : 0.00
REMARK 3
REMARK 3 ESTIMATED OVERALL COORDINATE ERROR.
REMARK 3 ESU BASED ON R VALUE (A) : 0.174
REMARK 3 ESU BASED ON FREE R VALUE (A) : 0.156
REMARK 3 ESU BASED ON MAXIMUM LIKELIHOOD (A) : 0.147
REMARK 3 ESU FOR B VALUES BASED ON MAXIMUM LIKELIHOOD (A**2) : 5.588
REMARK 3
REMARK 3 CORRELATION COEFFICIENTS.
REMARK 3 CORRELATION COEFFICIENT FO-FC : 0.961
REMARK 3 CORRELATION COEFFICIENT FO-FC FREE : 0.947
REMARK 3
REMARK 3 RMS DEVIATIONS FROM IDEAL VALUES COUNT RMS WEIGHT
REMARK 3 BOND LENGTHS REFINED (A) : 2973 ; 0.012 ; 0.021
REMARK 3 BOND LENGTHS OTHERS (A) : 2561 ; 0.001 ; 0.020
REMARK 3 BOND ANGLES REFINED (DEGREES) : 4044 ; 1.374 ; 1.949
REMARK 3 BOND ANGLES OTHERS (DEGREES) : 5979 ; 0.722 ; 3.000
REMARK 3 TORSION ANGLES, PERIOD 1 (DEGREES) : 352 ; 4.018 ; 3.000
REMARK 3 TORSION ANGLES, PERIOD 3 (DEGREES) : 515 ; 17.698 ; 15.000
REMARK 3 CHIRAL-CENTER RESTRAINTS (A**3) : 416 ; 0.086 ; 0.200
REMARK 3 GENERAL PLANES REFINED (A) : 3333 ; 0.005 ; 0.020
REMARK 3 GENERAL PLANES OTHERS (A) : 604 ; 0.002 ; 0.020
REMARK 3 NON-BONDED CONTACTS REFINED (A) : 714 ; 0.218 ; 0.300
REMARK 3 NON-BONDED CONTACTS OTHERS (A) : 2499 ; 0.204 ; 0.300
REMARK 3 H-BOND (X...Y) REFINED (A) : 259 ; 0.152 ; 0.500
REMARK 3 H-BOND (X...Y) OTHERS (A) : 4 ; 0.087 ; 0.500
REMARK 3 SYMMETRY VDW REFINED (A) : 18 ; 0.245 ; 0.300
REMARK 3 SYMMETRY VDW OTHERS (A) : 72 ; 0.248 ; 0.300

```

REMARK 3 SYMMETRY H-BOND REFINED (A): 13 ; 0.255 ; 0.500
 REMARK 3 SYMMETRY H-BOND OTHERS (A): 1 ; 0.052 ; 0.500
 REMARK 3
 REMARK 3 ISOTROPIC THERMAL FACTOR RESTRAINTS COUNT RMS WEIGHT
 REMARK 3 MAIN-CHAIN BOND REFINED (A**2): 1777 ; 0.618 ; 1.500
 REMARK 3 MAIN-CHAIN ANGLE REFINED (A**2): 2862 ; 1.177 ; 2.000
 REMARK 3 SIDE-CHAIN BOND REFINED (A**2): 1196 ; 1.812 ; 3.000
 REMARK 3 SIDE-CHAIN ANGLE REFINED (A**2): 1182 ; 3.002 ; 4.500
 REMARK 3
 REMARK 3 NCS RESTRAINTS STATISTICS
 REMARK 3 NUMBER OF NCS GROUPS : NULL
 REMARK 3
 REMARK 3 TLS DETAILS
 REMARK 3 NUMBER OF TLS GROUPS : 1
 REMARK 3
 REMARK 3 TLS GROUP : 1
 REMARK 3 NUMBER OF COMPONENTS GROUP : 2
 REMARK 3 COMPONENTS C SSSEQI TO C SSSEQI
 REMARK 3 RESIDUE RANGE : A 15 A 451
 REMARK 3 RESIDUE RANGE : S 795 S 823
 REMARK 3 ORIGIN FOR THE GROUP (A): 21.6620 27.4620 28.2370
 REMARK 3 T TENSOR
 REMARK 3 T11: 0.1474 T22: 0.0149
 REMARK 3 T33: 0.0919 T12: -0.0099
 REMARK 3 T13: -0.0455 T23: 0.0363
 REMARK 3 L TENSOR
 REMARK 3 L11: 1.0098 L22: 2.2577
 REMARK 3 L33: 1.2037 L12: 0.6963
 REMARK 3 L13: 0.4840 L23: 1.0420
 REMARK 3 S TENSOR
 REMARK 3 S11: 0.0288 S12: -0.1525 S13: -0.0400
 REMARK 3 S21: 0.1459 S22: 0.0002 S23: 0.1021
 REMARK 3 S31: 0.1876 S32: -0.0468 S33: -0.0290
 REMARK 3
 REMARK 3 BULK SOLVENT MODELLING.
 REMARK 3 METHOD USED : BABINET MODEL WITH MASK
 REMARK 3 PARAMETERS FOR MASK CALCULATION
 REMARK 3 VDW PROBE RADIUS : 1.40
 REMARK 3 ION PROBE RADIUS : 0.80
 REMARK 3 SHRINKAGE RADIUS : 0.80
 REMARK 3
 REMARK 3 OTHER REFINEMENT REMARKS: HYDROGENS HAVE BEEN ADDED IN THE
 REMARK 3 RIDING POSITIONS
 REMARK 4
 REMARK 4 1H2K COMPLIES WITH FORMAT V. 2.3, 09-JULY-1998
 REMARK 100
 REMARK 100 THIS ENTRY HAS BEEN PROCESSED BY EBI ON 12-AUG-2002.
 REMARK 100 THE EBI ID CODE IS EBI-11170.
 REMARK 200
 REMARK 200 EXPERIMENTAL DETAILS
 REMARK 200 EXPERIMENT TYPE : X-RAY DIFFRACTION
 REMARK 200 DATE OF DATA COLLECTION : 15-MAY-2002
 REMARK 200 TEMPERATURE (KELVIN) : 100
 REMARK 200 PH : 7.5
 REMARK 200 NUMBER OF CRYSTALS USED : 1
 REMARK 200
 REMARK 200 SYNCHROTRON (Y/N) : Y
 REMARK 200 RADIATION SOURCE : SRS BEAMLINE PX9.6
 REMARK 200 BEAMLINE : PX9.6
 REMARK 200 X-RAY GENERATOR MODEL : NULL
 REMARK 200 MONOCHROMATIC OR LAUE (M/L) : M

REMARK 200 WAVELENGTH OR RANGE (A) : 0.87
 REMARK 200 MONOCHROMATOR : NULL
 REMARK 200 OPTICS : NULL
 REMARK 200
 REMARK 200 DETECTOR TYPE : CCD
 REMARK 200 DETECTOR MANUFACTURER : ADSC
 REMARK 200 INTENSITY-INTEGRATION SOFTWARE : MOSFLM
 REMARK 200 DATA SCALING SOFTWARE : SCALA
 REMARK 200
 REMARK 200 NUMBER OF UNIQUE REFLECTIONS : 30574
 REMARK 200 RESOLUTION RANGE HIGH (A) : 2.15
 REMARK 200 RESOLUTION RANGE LOW (A) : 18.17
 REMARK 200 REJECTION CRITERIA (SIGMA(I)) : NONE
 REMARK 200
 REMARK 200 OVERALL.
 REMARK 200 COMPLETENESS FOR RANGE (%) : 99.2
 REMARK 200 DATA REDUNDANCY : 6.3
 REMARK 200 R MERGE (I) : 0.052
 REMARK 200 R SYM (I) : NULL
 REMARK 200 <I/SIGMA(I)> FOR THE DATA SET : 9.9
 REMARK 200
 REMARK 200 IN THE HIGHEST RESOLUTION SHELL.
 REMARK 200 HIGHEST RESOLUTION SHELL, RANGE HIGH (A) : 2.15
 REMARK 200 HIGHEST RESOLUTION SHELL, RANGE LOW (A) : 2.27
 REMARK 200 COMPLETENESS FOR SHELL (%) : 96.0
 REMARK 200 DATA REDUNDANCY IN SHELL : 3.4
 REMARK 200 R MERGE FOR SHELL (I) : 0.331
 REMARK 200 R SYM FOR SHELL (I) : NULL
 REMARK 200 <I/SIGMA(I)> FOR SHELL : 1.5
 REMARK 200
 REMARK 200 DIFFRACTION PROTOCOL: SINGLE WAVELENGTH
 REMARK 200 METHOD USED TO DETERMINE THE STRUCTURE: MAD
 REMARK 200 SOFTWARE USED: SOLVE
 REMARK 200 STARTING MODEL: NULL
 REMARK 200
 REMARK 200 REMARK: NULL
 REMARK 280
 REMARK 280 CRYSTAL
 REMARK 280 SOLVENT CONTENT, VS (%) : 63
 REMARK 280 MATTHEWS COEFFICIENT, VM (ANGSTROMS**3/DA) : 3.4
 REMARK 280
 REMARK 280 CRYSTALLIZATION CONDITIONS: 1.2M AMMONIUM SULPHATE, 4% PEG400,
 REMARK 280 0.1M HEPES PH7.5, ARGON ATMOSPHERE, 11MG/ML PROTEIN WITH
 REMARK 280 1MM FE(II), 2.5MM NOG AND 2.5MM PEPTIDE
 REMARK 290
 REMARK 290 CRYSTALLOGRAPHIC SYMMETRY
 REMARK 290 SYMMETRY OPERATORS FOR SPACE GROUP: P 41 21 2
 REMARK 290
 REMARK 290 SYMOP SYMMETRY
 REMARK 290 NNNMMM OPERATOR
 REMARK 290 1555 X, Y, Z
 REMARK 290 2555 -X, -Y, 1/2+Z
 REMARK 290 3555 1/2-Y, 1/2+X, 1/4+Z
 REMARK 290 4555 1/2+Y, 1/2-X, 3/4+Z
 REMARK 290 5555 1/2-X, 1/2+Y, 1/4-Z
 REMARK 290 6555 1/2+X, 1/2-Y, 3/4-Z
 REMARK 290 7555 Y, X, -Z
 REMARK 290 8555 -Y, -X, 1/2-Z
 REMARK 290
 REMARK 290 WHERE NNN -> OPERATOR NUMBER
 REMARK 290 MMM -> TRANSLATION VECTOR

REMARK 290

REMARK 290 CRYSTALLOGRAPHIC SYMMETRY TRANSFORMATIONS

REMARK 290 THE FOLLOWING TRANSFORMATIONS OPERATE ON THE ATOM/HETATM

REMARK 290 RECORDS IN THIS ENTRY TO PRODUCE CRYSTALLOGRAPHICALLY

REMARK 290 RELATED MOLECULES.

REMARK 290	SMTRY1	1	1.000000	0.000000	0.000000	0.000000
REMARK 290	SMTRY2	1	0.000000	1.000000	0.000000	0.000000
REMARK 290	SMTRY3	1	0.000000	0.000000	1.000000	0.000000
REMARK 290	SMTRY1	2	-1.000000	0.000000	0.000000	0.000000
REMARK 290	SMTRY2	2	0.000000	-1.000000	0.000000	0.000000
REMARK 290	SMTRY3	2	0.000000	0.000000	1.000000	73.32800
REMARK 290	SMTRY1	3	0.000000	-1.000000	0.000000	43.08050
REMARK 290	SMTRY2	3	1.000000	0.000000	0.000000	43.08050
REMARK 290	SMTRY3	3	0.000000	0.000000	1.000000	36.66400
REMARK 290	SMTRY1	4	0.000000	1.000000	0.000000	43.08050
REMARK 290	SMTRY2	4	-1.000000	0.000000	0.000000	43.08050
REMARK 290	SMTRY3	4	0.000000	0.000000	1.000000	109.99200
REMARK 290	SMTRY1	5	-1.000000	0.000000	0.000000	43.08050
REMARK 290	SMTRY2	5	0.000000	1.000000	0.000000	43.08050
REMARK 290	SMTRY3	5	0.000000	0.000000	-1.000000	36.66400
REMARK 290	SMTRY1	6	1.000000	0.000000	0.000000	43.08050
REMARK 290	SMTRY2	6	0.000000	-1.000000	0.000000	43.08050
REMARK 290	SMTRY3	6	0.000000	0.000000	-1.000000	109.99200
REMARK 290	SMTRY1	7	0.000000	1.000000	0.000000	0.000000
REMARK 290	SMTRY2	7	1.000000	0.000000	0.000000	0.000000
REMARK 290	SMTRY3	7	0.000000	0.000000	-1.000000	0.000000
REMARK 290	SMTRY1	8	0.000000	-1.000000	0.000000	0.000000
REMARK 290	SMTRY2	8	-1.000000	0.000000	0.000000	0.000000
REMARK 290	SMTRY3	8	0.000000	0.000000	-1.000000	73.32800

REMARK 290

REMARK 290 REMARK: NULL

REMARK 300

REMARK 300 BIOMOLECULE: 1

REMARK 300 THIS ENTRY CONTAINS THE CRYSTALLOGRAPHIC ASYMMETRIC UNIT

REMARK 300 WHICH CONSISTS OF 2 CHAIN(S). SEE REMARK 350 FOR

REMARK 300 INFORMATION ON GENERATING THE BIOLOGICAL MOLECULE(S).

REMARK 300

REMARK 300 QUATERNARY STRUCTURE FOR THIS ENTRY: TETRAMERIC

REMARK 300

REMARK 300 THE PROTEIN IS A HOMODIMER FORMED BY CHAIN A.

REMARK 300 A HETERODIMERIC ASSOCIATION OF CHAIN A WITH CHAIN S

REMARK 300 PRODUCES A TETRAMER.

REMARK 300

REMARK 300 THE BURIED SURFACE AREA SHOWN BELOW IS AN AVERAGE

REMARK 300 CALCULATED FOR THE HETEROTETRAMER AND DOES NOT

REMARK 300 CORRESPOND TO THE BURIED SURFACE AREA FOR THE

REMARK 300 HOMODIMER OF CHAIN A

REMARK 300

REMARK 300 THE HETERO-ASSEMBLY DESCRIBED BY REMARK 350 APPEARS

REMARK 300 TO BE A CASE OF STRONG CRYSTAL PACKING WITH

REMARK 300 THE MEAN DIFFERENCE IN ACCESSIBLE SURFACE AREA PER

REMARK 300 CHAIN BETWEEN THE ISOLATED CHAIN AND THAT FOR

REMARK 300 THE CHAIN IN THE COMPLEX IS 2203.4 ANGSTROM**2

REMARK 350

REMARK 350 GENERATING THE BIOMOLECULE

REMARK 350 COORDINATES FOR A COMPLETE MULTIMER REPRESENTING THE KNOWN

REMARK 350 BIOLOGICALLY SIGNIFICANT OLIGOMERIZATION STATE OF THE

REMARK 350 MOLECULE CAN BE GENERATED BY APPLYING BIOMT TRANSFORMATIONS

REMARK 350 GIVEN BELOW. BOTH NON-CRYSTALLOGRAPHIC AND

REMARK 350 CRYSTALLOGRAPHIC OPERATIONS ARE GIVEN.

REMARK 350

REMARK 350 BIOMOLECULE: 1
 REMARK 350 APPLY THE FOLLOWING TO CHAINS: A, S

REMARK 350	BIOMT1	1	1.000000	0.000000	0.000000	0.000000
REMARK 350	BIOMT2	1	0.000000	1.000000	0.000000	0.000000
REMARK 350	BIOMT3	1	0.000000	0.000000	1.000000	0.000000
REMARK 350	BIOMT1	2	0.000000	-1.000000	0.000000	86.16100
REMARK 350	BIOMT2	2	-1.000000	0.000000	0.000000	86.16100
REMARK 350	BIOMT3	2	0.000000	0.000000	-1.000000	73.32800

REMARK 465

REMARK 465 MISSING RESIDUES

REMARK 465 THE FOLLOWING RESIDUES WERE NOT LOCATED IN THE

REMARK 465 EXPERIMENT. (M=MODEL NUMBER; RES=RESIDUE NAME; C=CHAIN

REMARK 465 IDENTIFIER; SSSEQ=SEQUENCE NUMBER; I=INSERTION CODE.)

REMARK 465

REMARK 465 M RES C SSSEQI

REMARK 465	MET A	1
REMARK 465	ALA A	2
REMARK 465	ALA A	3
REMARK 465	THR A	4
REMARK 465	ALA A	5
REMARK 465	ALA A	6
REMARK 465	GLU A	7
REMARK 465	ALA A	8
REMARK 465	VAL A	9
REMARK 465	ALA A	10
REMARK 465	SER A	11
REMARK 465	GLY A	12
REMARK 465	SER A	13
REMARK 465	GLY A	14
REMARK 465	LYS A	304
REMARK 465	ARG A	305
REMARK 465	ILE A	306
REMARK 465	SER S	786
REMARK 465	MET S	787
REMARK 465	ASP S	788
REMARK 465	GLU S	789
REMARK 465	SER S	790
REMARK 465	GLY S	791
REMARK 465	LEU S	792
REMARK 465	PRO S	793
REMARK 465	GLN S	794
REMARK 465	GLN S	807
REMARK 465	GLY S	808
REMARK 465	SER S	809
REMARK 465	ARG S	810
REMARK 465	ASN S	811
REMARK 465	GLN S	824
REMARK 465	VAL S	825
REMARK 465	ASN S	826

REMARK 470

REMARK 470 MISSING ATOM

REMARK 470 THE FOLLOWING RESIDUES HAVE MISSING ATOMS (M=MODEL NUMBER;

REMARK 470 RES=RESIDUE NAME; C=CHAIN IDENTIFIER; SSEQ=SEQUENCE NUMBER;

REMARK 470 I=INSERTION CODE):

REMARK 470 M RES CSSEQI ATOMS

REMARK 470	GLU A	15	CG	CD	OE1	OE2
REMARK 470	GLU A	29	CG	CD	OE1	OE2
REMARK 470	ASN A	87	CG	OD1	ND2	
REMARK 470	LYS A	106	CD	CE	NZ	
REMARK 470	LYS A	115	CG	CD	CE	NZ
REMARK 470	ARG A	117	CG	CD	NE	CZ NH1 NH2

REMARK 470 GLN A 133 CG CD OE1 NE2
 REMARK 470 GLN A 136 CG CD OE1 NE2
 REMARK 470 GLN A 137 CG CD OE1 NE2
 REMARK 470 ARG A 156 CG CD NE CZ NH1 NH2
 REMARK 470 LYS A 157 CD CE NZ
 REMARK 470 LYS A 311 CG CD CE NZ
 REMARK 500
 REMARK 500 GEOMETRY AND STEREOCHEMISTRY
 REMARK 500 SUBTOPIC: COVALENT BOND ANGLES
 REMARK 500
 REMARK 500 THE STEREOCHEMICAL PARAMETERS OF THE FOLLOWING RESIDUES
 REMARK 500 HAVE VALUES WHICH DEVIATE FROM EXPECTED VALUES BY MORE
 REMARK 500 THAN 6*RMSD (M=MODEL NUMBER; RES=RESIDUE NAME; C=CHAIN
 REMARK 500 IDENTIFIER; SSEQ=SEQUENCE NUMBER; I=INSERTION CODE).
 REMARK 500
 REMARK 500 STANDARD TABLE:
 REMARK 500 FORMAT: (10X,I3,1X,A3,1X,A1,I4,A1,3(1X,A4,2X),12X,F5.1)
 REMARK 500
 REMARK 500 EXPECTED VALUES: ENGH AND HUBER, 1991
 REMARK 500
 REMARK 500 M RES CSSEQI ATM1 ATM2 ATM3
 REMARK 500 ASN A 84 N - CA - C ANGL. DEV. = 9.3 DEGREES
 REMARK 500
 REMARK 500 REMARK: NULL
 REMARK 500
 REMARK 500 GEOMETRY AND STEREOCHEMISTRY
 REMARK 500 SUBTOPIC: COVALENT BOND LENGTHS
 REMARK 500
 REMARK 500 THE STEREOCHEMICAL PARAMETERS OF THE FOLLOWING RESIDUES
 REMARK 500 HAVE VALUES WHICH DEVIATE FROM EXPECTED VALUES BY MORE
 REMARK 500 THAN 6*RMSD AND BY MORE THAN 0.150 ANGSTROMS (M=MODEL
 REMARK 500 NUMBER; RES=RESIDUE NAME; C=CHAIN IDENTIFIER; SSEQ=SEQUENCE
 REMARK 500 NUMBER; I=INSERTION CODE).
 REMARK 500
 REMARK 500 STANDARD TABLE:
 REMARK 500 FORMAT: (10X,I3,1X,A3,1X,A1,I4,A1,1X,2(A4,A1,3X),12X,F5.3)
 REMARK 500
 REMARK 500 EXPECTED VALUESS: ENGH AND HUBER, 1991
 REMARK 500
 REMARK 500 M RES CSSEQI ATM1 RES CSSEQI ATM2 DEVIATION
 REMARK 500 MET A 343 SD MET A 343 CE -0.249
 REMARK 500
 REMARK 500 REMARK: NULL
 REMARK 500
 REMARK 500 GEOMETRY AND STEREOCHEMISTRY
 REMARK 500 SUBTOPIC: CLOSE CONTACTS IN SAME ASYMMETRIC UNIT
 REMARK 500
 REMARK 500 THE FOLLOWING ATOMS ARE IN CLOSE CONTACT.
 REMARK 500

ATM1	RES	C	SSEQI	ATM2	RES	C	SSEQI	DISTANCE
O	GLN	A	209	O	HOH	Z	108	2.20

 REMARK 500
 REMARK 525
 REMARK 525 SOLVENT
 REMARK 525
 REMARK 525 THE SOLVENT MOLECULES ARE GIVEN CHAIN IDENTIFIERS TO
 REMARK 525 INDICATE THE PROTEIN CHAIN TO WHICH THEY ARE MOST CLOSELY
 REMARK 525 ASSOCIATED WITH:
 REMARK 525 PROTEIN CHAIN SOLVENT CHAIN
 REMARK 525 A Z
 REMARK 525 S H

REMARK 525
 REMARK 525 THE FOLLOWING SOLVENT MOLECULES LIE FARTHER THAN EXPECTED
 REMARK 525 FROM THE PROTEIN OR NUCLEIC ACID MOLECULE AND MAY BE
 REMARK 525 ASSOCIATED WITH A SYMMETRY RELATED MOLECULE (M=MODEL
 REMARK 525 NUMBER; RES=RESIDUE NAME; C=CHAIN IDENTIFIER; SSEQ=SEQUENCE
 REMARK 525 NUMBER; I=INSERTION CODE):
 REMARK 525
 REMARK 525 THESE MOLECULES CAN BE PLACED WITHIN 5.00 ANGSTROM OF THE
 REMARK 525 OBSERVED OLIGOMER BY APPLYING THE SYMMETRY TRANSFORMATION
 REMARK 525 INDICATED.
 REMARK 525

REMARK 525	M	RES	CSSEQI	ORIGINAL COORDINATES			SYMMETRY TRANS.		DIST.
REMARK 525				X	Y	Z			
REMARK 525	1	HOH	W 531	12.359	41.757	15.368	005	545	2.38
REMARK 525	1	HOH	W 609	10.971	45.216	18.991	005	545	3.25
REMARK 525	1	HOH	W 576	42.075	52.163	47.994	008	665	2.28
REMARK 525	1	HOH	W 687	28.879	5.577	12.106	005	555	2.82
REMARK 525	1	HOH	W 674	24.396	12.792	8.360	005	555	3.20
REMARK 525	1	HOH	W 543	27.797	7.178	14.664	005	555	2.83
REMARK 525	1	HOH	W 607	26.874	53.406	28.524	008	665	2.97

REMARK 600
 REMARK 600 HETEROGEN
 REMARK 600
 REMARK 600 FOR METAL ATOM FE FE2 A1350 THE COORDINATION ANGLES ARE:

REMARK 600	1	HIS	199A	NE2				
REMARK 600	2	ASP	201A	OD2	104.0			
REMARK 600	3	HIS	279A	NE2	85.8	88.0		
REMARK 600	4	OGA	1351A	O2	163.5	92.4	96.8	
REMARK 600	5	OGA	1351A	O2'	86.4	168.8	97.0	77.1
REMARK 600					1	2	3	4

REMARK 700
 REMARK 700 SHEET
 REMARK 700 THE SHEET STRUCTURE OF THIS MOLECULE IS BIFURCATED. IN
 REMARK 700 ORDER TO REPRESENT THIS FEATURE IN THE SHEET RECORDS BELOW,
 REMARK 700 TWO SHEETS ARE DEFINED.
 REMARK 800
 REMARK 800 SITE
 REMARK 800 SITE_IDENTIFIER: FE1
 REMARK 800 SITE_DESCRIPTION: FE BINDING SITE FOR CHAIN A
 REMARK 800
 REMARK 800 SITE_IDENTIFIER: OGA
 REMARK 800 SITE_DESCRIPTION: OGA BINDING SITE FOR CHAIN A
 REMARK 800
 REMARK 800 SITE_IDENTIFIER: SO1
 REMARK 800 SITE_DESCRIPTION: SO4 BINDING SITE FOR CHAIN A
 REMARK 800
 REMARK 800 SITE_IDENTIFIER: SO2
 REMARK 800 SITE_DESCRIPTION: SO4 BINDING SITE FOR CHAIN A
 REMARK 900
 REMARK 900 RELATED ENTRIES
 REMARK 900 RELATED ID: 1D7G RELATED DB: PDB
 REMARK 900 A MODEL FOR THE COMPLEX BETWEEN THE
 REMARK 900 HYPOXIA-INDUCIBLE FACTOR-1 (HIF-1) AND ITS
 REMARK 900 CONSENSUS DEOXYRIBONUCLEIC ACID SEQUENCE
 REMARK 900 RELATED ID: 1H2L RELATED DB: PDB
 REMARK 900 FACTOR INHIBITING HIF-1 ALPHA IN COMPLEX
 REMARK 900 WITH HIF-1 ALPHA FRAGMENT PEPTIDE
 REMARK 900 RELATED ID: 1H2M RELATED DB: PDB
 REMARK 900 FACTOR INHIBITING HIF-1 ALPHA IN COMPLEX
 REMARK 900 WITH HIF-1 ALPHA FRAGMENT PEPTIDE
 REMARK 900 RELATED ID: 1H2N RELATED DB: PDB

REMARK 900 FACTOR INHIBITING HIF-1 ALPHA IN COMPLEX
 REMARK 900 WITH HIF-1 ALPHA FRAGMENT PEPTIDE
 REMARK 900 RELATED ID: 1L8C RELATED DB: PDB
 REMARK 900 STRUCTURAL BASIS FOR HIF-1ALPHA/CBP
 REMARK 900 RECOGNITION IN THECELLULAR HYPOXIC RESPONSE
 REMARK 900 RELATED ID: 1LM8 RELATED DB: PDB
 REMARK 900 STRUCTURE OF A HIF-1A-PVHL-ELONGINB-
 REMARK 900 ELONGINC COMPLEX
 REMARK 900 RELATED ID: 1LQB RELATED DB: PDB
 REMARK 900 CRYSTAL STRUCTURE OF A HYDROXYLATED HIF-1
 REMARK 900 ALPHA PEPTIDEBOUND TO THE PVHL/ELONGIN-C/
 REMARK 900 ELONGIN-B COMPLEX

DBREF 1H2K A 1 349 SWS Q969Q7 Q969Q7 1 349
 DBREF 1H2K S 786 826 SWS Q16665 HIFA_HUMAN 786 826
 SEQRES 1 A 349 MET ALA ALA THR ALA ALA GLU ALA VAL ALA SER GLY SER
 SEQRES 2 A 349 GLY GLU PRO ARG GLU GLU ALA GLY ALA LEU GLY PRO ALA
 SEQRES 3 A 349 TRP ASP GLU SER GLN LEU ARG SER TYR SER PHE PRO THR
 SEQRES 4 A 349 ARG PRO ILE PRO ARG LEU SER GLN SER ASP PRO ARG ALA
 SEQRES 5 A 349 GLU GLU LEU ILE GLU ASN GLU GLU PRO VAL VAL LEU THR
 SEQRES 6 A 349 ASP THR ASN LEU VAL TYR PRO ALA LEU LYS TRP ASP LEU
 SEQRES 7 A 349 GLU TYR LEU GLN GLU ASN ILE GLY ASN GLY ASP PHE SER
 SEQRES 8 A 349 VAL TYR SER ALA SER THR HIS LYS PHE LEU TYR TYR ASP
 SEQRES 9 A 349 GLU LYS LYS MET ALA ASN PHE GLN ASN PHE LYS PRO ARG
 SEQRES 10 A 349 SER ASN ARG GLU GLU MET LYS PHE HIS GLU PHE VAL GLU
 SEQRES 11 A 349 LYS LEU GLN ASP ILE GLN GLN ARG GLY GLY GLU GLU ARG
 SEQRES 12 A 349 LEU TYR LEU GLN GLN THR LEU ASN ASP THR VAL GLY ARG
 SEQRES 13 A 349 LYS ILE VAL MET ASP PHE LEU GLY PHE ASN TRP ASN TRP
 SEQRES 14 A 349 ILE ASN LYS GLN GLN GLY LYS ARG GLY TRP GLY GLN LEU
 SEQRES 15 A 349 THR SER ASN LEU LEU LEU ILE GLY MET GLU GLY ASN VAL
 SEQRES 16 A 349 THR PRO ALA HIS TYR ASP GLU GLN GLN ASN PHE PHE ALA
 SEQRES 17 A 349 GLN ILE LYS GLY TYR LYS ARG CYS ILE LEU PHE PRO PRO
 SEQRES 18 A 349 ASP GLN PHE GLU CYS LEU TYR PRO TYR PRO VAL HIS HIS
 SEQRES 19 A 349 PRO CYS ASP ARG GLN SER GLN VAL ASP PHE ASP ASN PRO
 SEQRES 20 A 349 ASP TYR GLU ARG PHE PRO ASN PHE GLN ASN VAL VAL GLY
 SEQRES 21 A 349 TYR GLU THR VAL VAL GLY PRO GLY ASP VAL LEU TYR ILE
 SEQRES 22 A 349 PRO MET TYR TRP TRP HIS HIS ILE GLU SER LEU LEU ASN
 SEQRES 23 A 349 GLY GLY ILE THR ILE THR VAL ASN PHE TRP TYR LYS GLY
 SEQRES 24 A 349 ALA PRO THR PRO LYS ARG ILE GLU TYR PRO LEU LYS ALA
 SEQRES 25 A 349 HIS GLN LYS VAL ALA ILE MET ARG ASN ILE GLU LYS MET
 SEQRES 26 A 349 LEU GLY GLU ALA LEU GLY ASN PRO GLN GLU VAL GLY PRO
 SEQRES 27 A 349 LEU LEU ASN THR MET ILE LYS GLY ARG TYR ASN
 SEQRES 1 S 41 SER MET ASP GLU SER GLY LEU PRO GLN LEU THR SER TYR
 SEQRES 2 S 41 ASP CYS GLU VAL ASN ALA PRO ILE GLN GLY SER ARG ASN
 SEQRES 3 S 41 LEU LEU GLN GLY GLU GLU LEU LEU ARG ALA LEU ASP GLN
 SEQRES 4 S 41 VAL ASN
 HET FE2 A1350 1
 HET OGA A1351 10
 HET SO4 A1352 5
 HET SO4 A1353 5
 HETNAM FE2 FE (II) ION
 HETNAM OGA N-OXALYOLGLYCINE
 HETNAM SO4 SULFATE ION
 FORMUL 3 FE2 FE1 2+
 FORMUL 4 OGA C4 H5 N1 O5
 FORMUL 5 SO4 2(O4 S1 2-)
 FORMUL 6 HOH *194(H2 O1)
 HELIX 1 1 ASP A 28 LEU A 32 5
 HELIX 2 2 ASP A 49 ASN A 58 1
 HELIX 3 3 VAL A 70 TRP A 76 5
 HELIX 4 4 ASP A 77 ILE A 85 1
 HELIX 5 5 ASP A 104 GLN A 112 5

5
 10
 7
 9
 9

[illegible]

ATOM	12	CD	PRO	A	16	5.108	30.911	11.548	1.00	50.56	C
ATOM	13	N	ARG	A	17	7.133	26.746	10.550	1.00	48.83	N
ATOM	14	CA	ARG	A	17	7.219	25.695	9.550	1.00	48.33	C
ATOM	15	C	ARG	A	17	5.967	24.832	9.561	1.00	47.48	C
ATOM	16	O	ARG	A	17	5.245	24.782	10.557	1.00	47.64	O
ATOM	17	CB	ARG	A	17	8.421	24.798	9.835	1.00	48.61	C
ATOM	18	CG	ARG	A	17	9.776	25.511	9.835	1.00	49.21	C
ATOM	19	CD	ARG	A	17	10.944	24.577	10.196	1.00	49.84	C
ATOM	20	NE	ARG	A	17	10.918	24.137	11.596	1.00	50.12	N
ATOM	21	CZ	ARG	A	17	11.455	24.809	12.623	1.00	50.99	C
ATOM	22	NH1	ARG	A	17	12.065	25.979	12.431	1.00	50.40	N
ATOM	23	NH2	ARG	A	17	11.381	24.310	13.857	1.00	50.81	N
ATOM	24	N	GLU	A	18	5.723	24.153	8.446	1.00	46.05	N
ATOM	25	CA	GLU	A	18	4.603	23.246	8.329	1.00	45.06	C
ATOM	26	C	GLU	A	18	5.096	21.830	8.607	1.00	44.00	C
ATOM	27	O	GLU	A	18	6.101	21.405	8.044	1.00	43.89	O
ATOM	28	CB	GLU	A	18	4.013	23.324	6.923	1.00	45.21	C
ATOM	29	CG	GLU	A	18	3.323	24.648	6.621	1.00	45.56	C
ATOM	30	CD	GLU	A	18	1.951	24.780	7.265	1.00	45.64	C
ATOM	31	OE1	GLU	A	18	1.342	23.749	7.641	1.00	44.14	O
ATOM	32	OE2	GLU	A	18	1.480	25.932	7.388	1.00	45.83	O
ATOM	33	N	GLU	A	19	4.396	21.113	9.484	1.00	42.49	N
ATOM	34	CA	GLU	A	19	4.734	19.728	9.795	1.00	41.55	C
ATOM	35	C	GLU	A	19	4.357	18.817	8.635	1.00	39.79	C
ATOM	36	O	GLU	A	19	3.266	18.933	8.066	1.00	39.13	O
ATOM	37	CB	GLU	A	19	4.010	19.256	11.052	1.00	41.88	C
ATOM	38	CG	GLU	A	19	4.420	19.997	12.311	1.00	44.80	C
ATOM	39	CD	GLU	A	19	4.276	19.155	13.574	1.00	49.28	C
ATOM	40	OE1	GLU	A	19	3.759	18.008	13.497	1.00	51.52	O
ATOM	41	OE2	GLU	A	19	4.695	19.643	14.656	1.00	52.76	O
ATOM	42	N	ALA	A	20	5.270	17.910	8.311	1.00	38.00	N
ATOM	43	CA	ALA	A	20	5.099	16.952	7.227	1.00	36.77	C
ATOM	44	C	ALA	A	20	3.803	16.168	7.373	1.00	35.56	C
ATOM	45	O	ALA	A	20	3.445	15.734	8.460	1.00	35.74	O
ATOM	46	CB	ALA	A	20	6.283	15.999	7.180	1.00	36.63	C
ATOM	47	N	GLY	A	21	3.082	16.020	6.279	1.00	34.05	N
ATOM	48	CA	GLY	A	21	1.860	15.242	6.307	1.00	33.24	C
ATOM	49	C	GLY	A	21	0.666	16.137	6.551	1.00	32.36	C
ATOM	50	O	GLY	A	21	-0.393	15.673	6.951	1.00	30.93	O
ATOM	51	N	ALA	A	22	0.867	17.432	6.323	1.00	32.41	N
ATOM	52	CA	ALA	A	22	-0.184	18.425	6.459	1.00	32.78	C
ATOM	53	C	ALA	A	22	-0.723	18.441	7.873	1.00	33.14	C
ATOM	54	O	ALA	A	22	-1.915	18.605	8.088	1.00	32.74	O
ATOM	55	CB	ALA	A	22	-1.304	18.139	5.462	1.00	32.61	C
ATOM	56	N	LEU	A	23	0.151	18.253	8.849	1.00	34.01	N
ATOM	57	CA	LEU	A	23	-0.297	18.275	10.232	1.00	34.91	C
ATOM	58	C	LEU	A	23	-0.342	19.694	10.757	1.00	35.27	C
ATOM	59	O	LEU	A	23	-0.528	19.918	11.943	1.00	35.72	O
ATOM	60	CB	LEU	A	23	0.565	17.366	11.097	1.00	35.27	C
ATOM	61	CG	LEU	A	23	0.384	15.910	10.653	1.00	36.36	C
ATOM	62	CD1	LEU	A	23	1.211	14.947	11.491	1.00	37.75	C
ATOM	63	CD2	LEU	A	23	-1.077	15.523	10.719	1.00	37.98	C
ATOM	64	N	GLY	A	24	-0.177	20.656	9.855	1.00	35.73	N
ATOM	65	CA	GLY	A	24	-0.332	22.053	10.194	1.00	35.66	C
ATOM	66	C	GLY	A	24	0.901	22.655	10.804	1.00	36.08	C
ATOM	67	O	GLY	A	24	1.945	22.001	10.913	1.00	36.05	O
ATOM	68	N	PRO	A	25	0.764	23.894	11.253	1.00	35.96	N
ATOM	69	CA	PRO	A	25	1.896	24.628	11.804	1.00	36.12	C
ATOM	70	C	PRO	A	25	2.327	23.980	13.108	1.00	36.32	C
ATOM	71	O	PRO	A	25	1.488	23.577	13.914	1.00	36.08	O
ATOM	72	CB	PRO	A	25	1.341	26.047	12.043	1.00	36.15	C

ATOM	73	CG	PRO	A	25	-0.162	25.942	11.982	1.00	35.84	C
ATOM	74	CD	PRO	A	25	-0.504	24.638	11.350	1.00	36.22	C
ATOM	75	N	ALA	A	26	3.632	23.848	13.285	1.00	36.64	N
ATOM	76	CA	ALA	A	26	4.178	23.274	14.499	1.00	37.19	C
ATOM	77	C	ALA	A	26	3.860	24.172	15.706	1.00	36.77	C
ATOM	78	O	ALA	A	26	3.595	23.678	16.808	1.00	37.19	O
ATOM	79	CB	ALA	A	26	5.672	23.099	14.347	1.00	37.52	C
ATOM	80	N	TRP	A	27	3.848	25.479	15.484	1.00	35.62	N
ATOM	81	CA	TRP	A	27	3.520	26.420	16.543	1.00	35.04	C
ATOM	82	C	TRP	A	27	3.029	27.729	15.933	1.00	34.20	C
ATOM	83	O	TRP	A	27	2.992	27.883	14.723	1.00	33.57	O
ATOM	84	CB	TRP	A	27	4.774	26.672	17.382	1.00	35.28	C
ATOM	85	CG	TRP	A	27	5.951	26.889	16.511	1.00	34.67	C
ATOM	86	CD1	TRP	A	27	6.761	25.930	15.955	1.00	35.52	C
ATOM	87	CD2	TRP	A	27	6.426	28.135	16.033	1.00	34.40	C
ATOM	88	NE1	TRP	A	27	7.723	26.522	15.172	1.00	35.09	N
ATOM	89	CE2	TRP	A	27	7.541	27.877	15.209	1.00	34.23	C
ATOM	90	CE3	TRP	A	27	6.038	29.452	16.232	1.00	34.31	C
ATOM	91	CZ2	TRP	A	27	8.255	28.879	14.592	1.00	35.79	C
ATOM	92	CZ3	TRP	A	27	6.750	30.442	15.629	1.00	36.35	C
ATOM	93	CH2	TRP	A	27	7.847	30.154	14.808	1.00	36.47	C
ATOM	94	N	ASP	A	28	2.638	28.672	16.766	1.00	33.77	N
ATOM	95	CA	ASP	A	28	2.259	29.970	16.249	1.00	33.48	C
ATOM	96	C	ASP	A	28	2.759	31.050	17.165	1.00	32.26	C
ATOM	97	O	ASP	A	28	3.210	30.791	18.276	1.00	32.05	O
ATOM	98	CB	ASP	A	28	0.749	30.080	16.037	1.00	34.30	C
ATOM	99	CG	ASP	A	28	-0.026	29.986	17.317	1.00	36.29	C
ATOM	100	OD1	ASP	A	28	-0.184	31.027	18.002	1.00	39.59	O
ATOM	101	OD2	ASP	A	28	-0.517	28.906	17.712	1.00	39.78	O
ATOM	102	N	GLU	A	29	2.691	32.268	16.654	1.00	31.30	N
ATOM	103	CA	GLU	A	29	3.181	33.465	17.326	1.00	30.15	C
ATOM	104	C	GLU	A	29	2.674	33.625	18.752	1.00	28.85	C
ATOM	105	O	GLU	A	29	3.407	34.036	19.621	1.00	28.58	O
ATOM	106	CB	GLU	A	29	2.791	34.682	16.503	1.00	30.32	C
ATOM	107	N	SER	A	30	1.414	33.313	18.992	1.00	27.95	N
ATOM	108	CA	SER	A	30	0.845	33.501	20.320	1.00	27.46	C
ATOM	109	C	SER	A	30	1.537	32.671	21.389	1.00	26.93	C
ATOM	110	O	SER	A	30	1.312	32.907	22.567	1.00	26.79	O
ATOM	111	CB	SER	A	30	-0.651	33.168	20.322	1.00	27.23	C
ATOM	112	OG	SER	A	30	-0.857	31.764	20.306	1.00	27.69	O
ATOM	113	N	GLN	A	31	2.360	31.703	20.984	1.00	26.55	N
ATOM	114	CA	GLN	A	31	3.071	30.837	21.926	1.00	26.63	C
ATOM	115	C	GLN	A	31	4.419	31.409	22.334	1.00	26.66	C
ATOM	116	O	GLN	A	31	5.078	30.855	23.205	1.00	26.45	O
ATOM	117	CB	GLN	A	31	3.282	29.426	21.349	1.00	26.41	C
ATOM	118	CG	GLN	A	31	1.998	28.637	21.131	1.00	26.28	C
ATOM	119	CD	GLN	A	31	2.245	27.287	20.489	1.00	26.25	C
ATOM	120	OE1	GLN	A	31	2.258	27.183	19.271	1.00	27.92	O
ATOM	121	NE2	GLN	A	31	2.465	26.258	21.305	1.00	24.36	N
ATOM	122	N	LEU	A	32	4.824	32.508	21.703	1.00	26.89	N
ATOM	123	CA	LEU	A	32	6.083	33.176	22.029	1.00	27.24	C
ATOM	124	C	LEU	A	32	5.852	34.321	23.006	1.00	26.92	C
ATOM	125	O	LEU	A	32	4.888	35.047	22.868	1.00	26.42	O
ATOM	126	CB	LEU	A	32	6.717	33.746	20.760	1.00	27.46	C
ATOM	127	CG	LEU	A	32	6.964	32.728	19.640	1.00	28.65	C
ATOM	128	CD1	LEU	A	32	7.630	33.391	18.452	1.00	29.44	C
ATOM	129	CD2	LEU	A	32	7.792	31.573	20.127	1.00	27.98	C
ATOM	130	N	ARG	A	33	6.728	34.472	23.995	1.00	26.71	N
ATOM	131	CA	ARG	A	33	6.627	35.596	24.923	1.00	26.77	C
ATOM	132	C	ARG	A	33	7.040	36.880	24.209	1.00	26.57	C
ATOM	133	O	ARG	A	33	7.719	36.844	23.203	1.00	26.30	O

ATOM	134	CB	ARG	A	33	7.492	35.357	26.163	1.00	26.67	C
ATOM	135	CG	ARG	A	33	7.052	34.141	26.983	1.00	26.29	C
ATOM	136	CD	ARG	A	33	7.937	33.837	28.181	1.00	25.85	C
ATOM	137	NE	ARG	A	33	7.381	32.778	29.018	1.00	25.80	N
ATOM	138	CZ	ARG	A	33	6.451	32.946	29.945	1.00	24.68	C
ATOM	139	NH1	ARG	A	33	5.937	34.140	30.189	1.00	23.50	N
ATOM	140	NH2	ARG	A	33	6.029	31.901	30.637	1.00	25.35	N
ATOM	141	N	SER	A	34	6.633	38.020	24.732	1.00	26.80	N
ATOM	142	CA	SER	A	34	6.903	39.280	24.061	1.00	27.15	C
ATOM	143	C	SER	A	34	7.990	40.048	24.791	1.00	26.39	C
ATOM	144	O	SER	A	34	7.964	40.140	26.005	1.00	25.87	O
ATOM	145	CB	SER	A	34	5.628	40.104	24.030	1.00	27.68	C
ATOM	146	OG	SER	A	34	5.494	40.737	25.285	1.00	32.47	O
ATOM	147	N	TYR	A	35	8.944	40.589	24.042	1.00	25.89	N
ATOM	148	CA	TYR	A	35	10.110	41.223	24.637	1.00	26.28	C
ATOM	149	C	TYR	A	35	10.353	42.558	23.970	1.00	26.85	C
ATOM	150	O	TYR	A	35	9.722	42.856	22.967	1.00	26.92	O
ATOM	151	CB	TYR	A	35	11.326	40.308	24.510	1.00	25.90	C
ATOM	152	CG	TYR	A	35	11.169	39.032	25.309	1.00	25.33	C
ATOM	153	CD1	TYR	A	35	10.975	39.073	26.685	1.00	24.99	C
ATOM	154	CD2	TYR	A	35	11.198	37.787	24.692	1.00	23.82	C
ATOM	155	CE1	TYR	A	35	10.823	37.898	27.429	1.00	24.56	C
ATOM	156	CE2	TYR	A	35	11.046	36.620	25.426	1.00	24.42	C
ATOM	157	CZ	TYR	A	35	10.862	36.682	26.793	1.00	23.48	C
ATOM	158	OH	TYR	A	35	10.696	35.524	27.513	1.00	24.75	O
ATOM	159	N	SER	A	36	11.304	43.327	24.496	1.00	27.27	N
ATOM	160	CA	SER	A	36	11.525	44.704	24.052	1.00	27.55	C
ATOM	161	C	SER	A	36	12.513	44.912	22.917	1.00	27.06	C
ATOM	162	O	SER	A	36	12.734	46.049	22.504	1.00	27.94	O
ATOM	163	CB	SER	A	36	12.082	45.498	25.226	1.00	27.88	C
ATOM	164	OG	SER	A	36	13.350	44.976	25.590	1.00	28.36	O
ATOM	165	N	PHE	A	37	13.128	43.851	22.429	1.00	25.35	N
ATOM	166	CA	PHE	A	37	14.202	44.014	21.461	1.00	24.62	C
ATOM	167	C	PHE	A	37	13.899	43.272	20.159	1.00	24.75	C
ATOM	168	O	PHE	A	37	13.130	42.335	20.135	1.00	24.25	O
ATOM	169	CB	PHE	A	37	15.487	43.462	22.071	1.00	23.87	C
ATOM	170	CG	PHE	A	37	15.318	42.069	22.635	1.00	22.53	C
ATOM	171	CD1	PHE	A	37	15.348	40.972	21.802	1.00	21.11	C
ATOM	172	CD2	PHE	A	37	15.069	41.872	23.988	1.00	21.63	C
ATOM	173	CE1	PHE	A	37	15.158	39.687	22.314	1.00	22.41	C
ATOM	174	CE2	PHE	A	37	14.900	40.612	24.505	1.00	22.29	C
ATOM	175	CZ	PHE	A	37	14.936	39.509	23.675	1.00	21.45	C
ATOM	176	N	PRO	A	38	14.489	43.715	19.067	1.00	24.94	N
ATOM	177	CA	PRO	A	38	14.322	43.017	17.793	1.00	24.67	C
ATOM	178	C	PRO	A	38	15.267	41.823	17.678	1.00	24.41	C
ATOM	179	O	PRO	A	38	16.249	41.745	18.427	1.00	23.73	O
ATOM	180	CB	PRO	A	38	14.725	44.072	16.783	1.00	24.75	C
ATOM	181	CG	PRO	A	38	15.791	44.872	17.530	1.00	26.09	C
ATOM	182	CD	PRO	A	38	15.287	44.950	18.941	1.00	25.27	C
ATOM	183	N	THR	A	39	14.981	40.927	16.734	1.00	23.30	N
ATOM	184	CA	THR	A	39	15.859	39.816	16.444	1.00	23.42	C
ATOM	185	C	THR	A	39	15.857	39.534	14.955	1.00	24.29	C
ATOM	186	O	THR	A	39	14.958	39.964	14.239	1.00	24.61	O
ATOM	187	CB	THR	A	39	15.368	38.538	17.135	1.00	23.09	C
ATOM	188	OG1	THR	A	39	14.044	38.232	16.680	1.00	19.92	O
ATOM	189	CG2	THR	A	39	15.213	38.731	18.641	1.00	23.03	C
ATOM	190	N	ARG	A	40	16.854	38.773	14.525	1.00	24.43	N
ATOM	191	CA	ARG	A	40	16.982	38.273	13.170	1.00	24.91	C
ATOM	192	C	ARG	A	40	17.061	36.751	13.268	1.00	24.49	C
ATOM	193	O	ARG	A	40	17.434	36.225	14.301	1.00	23.58	O
ATOM	194	CB	ARG	A	40	18.253	38.805	12.536	1.00	25.36	C

ATOM	195	CG	ARG	A	40	18.208	40.281	12.248	1.00	30.74	
ATOM	196	CD	ARG	A	40	16.823	40.775	11.890	1.00	34.86	C
ATOM	197	NE	ARG	A	40	16.604	40.989	10.477	1.00	39.38	C
ATOM	198	CZ	ARG	A	40	15.403	41.229	9.971	1.00	43.86	N
ATOM	199	NH1	ARG	A	40	14.345	41.254	10.782	1.00	45.88	N
ATOM	200	NH2	ARG	A	40	15.252	41.457	8.674	1.00	44.96	N
ATOM	201	N	PRO	A	41	16.687	36.031	12.221	1.00	25.08	N
ATOM	202	CA	PRO	A	41	16.715	34.565	12.290	1.00	25.39	C
ATOM	203	C	PRO	A	41	18.095	33.913	12.282	1.00	25.10	C
ATOM	204	O	PRO	A	41	19.007	34.370	11.636	1.00	25.16	O
ATOM	205	CB	PRO	A	41	15.953	34.130	11.023	1.00	25.76	C
ATOM	206	CG	PRO	A	41	15.286	35.392	10.504	1.00	26.54	C
ATOM	207	CD	PRO	A	41	16.151	36.524	10.939	1.00	24.85	C
ATOM	208	N	ILE	A	42	18.225	32.823	13.020	1.00	25.28	N
ATOM	209	CA	ILE	A	42	19.388	31.974	12.919	1.00	24.31	C
ATOM	210	C	ILE	A	42	19.147	31.116	11.677	1.00	24.68	C
ATOM	211	O	ILE	A	42	18.043	30.614	11.466	1.00	24.72	O
ATOM	212	CB	ILE	A	42	19.481	31.104	14.163	1.00	24.80	C
ATOM	213	CG1	ILE	A	42	19.763	31.993	15.384	1.00	24.24	C
ATOM	214	CG2	ILE	A	42	20.530	30.004	13.961	1.00	23.81	C
ATOM	215	CD1	ILE	A	42	19.531	31.325	16.729	1.00	23.59	C
ATOM	216	N	PRO	A	43	20.146	30.953	10.826	1.00	24.48	N
ATOM	217	CA	PRO	A	43	19.963	30.108	9.651	1.00	24.60	C
ATOM	218	C	PRO	A	43	19.611	28.650	10.001	1.00	24.85	C
ATOM	219	O	PRO	A	43	20.148	28.130	10.989	1.00	24.38	O
ATOM	220	CB	PRO	A	43	21.320	30.192	8.937	1.00	24.87	C
ATOM	221	CG	PRO	A	43	22.040	31.372	9.541	1.00	25.04	C
ATOM	222	CD	PRO	A	43	21.475	31.583	10.886	1.00	24.37	C
ATOM	223	N	ARG	A	44	18.686	28.032	9.248	1.00	24.70	N
ATOM	224	CA	ARG	A	44	18.367	26.608	9.391	1.00	25.53	C
ATOM	225	C	ARG	A	44	18.910	25.943	8.152	1.00	25.00	C
ATOM	226	O	ARG	A	44	18.505	26.265	7.030	1.00	24.62	O
ATOM	227	CB	ARG	A	44	16.873	26.287	9.452	1.00	26.51	C
ATOM	228	CG	ARG	A	44	16.044	27.133	10.378	1.00	29.49	C
ATOM	229	CD	ARG	A	44	14.683	26.485	10.813	1.00	31.03	C
ATOM	230	NE	ARG	A	44	14.401	25.120	10.323	1.00	32.36	N
ATOM	231	CZ	ARG	A	44	14.174	24.057	11.126	1.00	33.85	C
ATOM	232	NH1	ARG	A	44	14.239	24.170	12.451	1.00	30.71	N
ATOM	233	NH2	ARG	A	44	13.898	22.863	10.613	1.00	35.56	N
ATOM	234	N	LEU	A	45	19.815	25.006	8.337	1.00	23.91	N
ATOM	235	CA	LEU	A	45	20.500	24.444	7.202	1.00	23.40	C
ATOM	236	C	LEU	A	45	20.684	22.967	7.352	1.00	23.52	C
ATOM	237	O	LEU	A	45	20.559	22.423	8.446	1.00	22.50	O
ATOM	238	CB	LEU	A	45	21.888	25.064	7.093	1.00	22.62	C
ATOM	239	CG	LEU	A	45	21.911	26.563	6.819	1.00	23.96	C
ATOM	240	CD1	LEU	A	45	23.317	27.111	6.947	1.00	24.52	C
ATOM	241	CD2	LEU	A	45	21.366	26.845	5.423	1.00	24.63	C
ATOM	242	N	SER	A	46	21.018	22.347	6.227	1.00	23.57	N
ATOM	243	CA	SER	A	46	21.382	20.975	6.221	1.00	24.12	C
ATOM	244	C	SER	A	46	22.820	20.888	6.668	1.00	24.81	C
ATOM	245	O	SER	A	46	23.640	21.757	6.398	1.00	23.76	O
ATOM	246	CB	SER	A	46	21.236	20.354	4.830	1.00	23.80	C
ATOM	247	OG	SER	A	46	21.744	19.020	4.830	1.00	24.09	O
ATOM	248	N	GLN	A	47	23.089	19.804	7.366	1.00	25.92	N
ATOM	249	CA	GLN	A	47	24.399	19.444	7.844	1.00	27.25	C
ATOM	250	C	GLN	A	47	25.379	19.326	6.674	1.00	27.54	C
ATOM	251	O	GLN	A	47	26.563	19.564	6.836	1.00	27.97	O
ATOM	252	CB	GLN	A	47	24.245	18.088	8.554	1.00	28.46	C
ATOM	253	CG	GLN	A	47	25.487	17.279	8.705	1.00	31.10	C
ATOM	254	CD	GLN	A	47	25.776	16.372	7.570	1.00	33.17	C
ATOM	255	OE1	GLN	A	47	24.881	15.970	6.803	1.00	36.23	O

ATOM	256	NE2	GLN	A	47	27.041	16.000	7.458	1.00	36.17	N
ATOM	257	N	SER	A	48	24.884	18.951	5.498	1.00	27.72	N
ATOM	258	CA	SER	A	48	25.736	18.809	4.323	1.00	28.35	C
ATOM	259	C	SER	A	48	26.016	20.150	3.629	1.00	28.86	C
ATOM	260	O	SER	A	48	26.825	20.235	2.711	1.00	28.96	O
ATOM	261	CB	SER	A	48	25.092	17.839	3.324	1.00	28.65	C
ATOM	262	OG	SER	A	48	23.798	18.276	2.918	1.00	28.51	O
ATOM	263	N	ASP	A	49	25.347	21.203	4.065	1.00	29.16	N
ATOM	264	CA	ASP	A	49	25.515	22.496	3.442	1.00	29.28	C
ATOM	265	C	ASP	A	49	26.800	23.163	3.913	1.00	29.71	C
ATOM	266	O	ASP	A	49	26.981	23.398	5.101	1.00	29.02	O
ATOM	267	CB	ASP	A	49	24.325	23.369	3.783	1.00	29.35	C
ATOM	268	CG	ASP	A	49	24.316	24.667	3.010	1.00	29.93	C
ATOM	269	OD1	ASP	A	49	25.398	25.183	2.666	1.00	28.14	O
ATOM	270	OD2	ASP	A	49	23.259	25.247	2.731	1.00	30.49	O
ATOM	271	N	PRO	A	50	27.693	23.483	2.976	1.00	30.34	N
ATOM	272	CA	PRO	A	50	28.964	24.134	3.315	1.00	30.54	C
ATOM	273	C	PRO	A	50	28.783	25.354	4.186	1.00	30.38	C
ATOM	274	O	PRO	A	50	29.661	25.651	4.990	1.00	30.53	O
ATOM	275	CB	PRO	A	50	29.520	24.573	1.952	1.00	30.39	C
ATOM	276	CG	PRO	A	50	28.926	23.667	0.988	1.00	30.96	C
ATOM	277	CD	PRO	A	50	27.574	23.243	1.531	1.00	30.51	C
ATOM	278	N	ARG	A	51	27.683	26.072	4.008	1.00	30.50	N
ATOM	279	CA	ARG	A	51	27.439	27.253	4.817	1.00	30.94	C
ATOM	280	C	ARG	A	51	27.341	26.875	6.299	1.00	30.49	C
ATOM	281	O	ARG	A	51	27.744	27.646	7.161	1.00	29.74	O
ATOM	282	CB	ARG	A	51	26.171	27.979	4.370	1.00	30.97	C
ATOM	283	CG	ARG	A	51	26.337	28.781	3.099	1.00	33.47	C
ATOM	284	CD	ARG	A	51	25.029	29.321	2.521	1.00	34.80	C
ATOM	285	NE	ARG	A	51	24.071	28.252	2.221	1.00	36.68	N
ATOM	286	CZ	ARG	A	51	22.766	28.448	2.081	1.00	37.94	C
ATOM	287	NH1	ARG	A	51	22.260	29.677	2.208	1.00	39.10	N
ATOM	288	NH2	ARG	A	51	21.967	27.436	1.794	1.00	36.51	N
ATOM	289	N	ALA	A	52	26.831	25.684	6.596	1.00	30.03	N
ATOM	290	CA	ALA	A	52	26.697	25.288	7.993	1.00	29.96	C
ATOM	291	C	ALA	A	52	28.079	25.101	8.593	1.00	30.07	C
ATOM	292	O	ALA	A	52	28.345	25.518	9.710	1.00	29.00	O
ATOM	293	CB	ALA	A	52	25.901	24.028	8.119	1.00	29.81	C
ATOM	294	N	GLU	A	53	28.958	24.455	7.845	1.00	30.12	N
ATOM	295	CA	GLU	A	53	30.290	24.213	8.347	1.00	31.08	C
ATOM	296	C	GLU	A	53	30.999	25.550	8.579	1.00	30.37	C
ATOM	297	O	GLU	A	53	31.683	25.730	9.580	1.00	29.89	O
ATOM	298	CB	GLU	A	53	31.086	23.362	7.379	1.00	31.41	C
ATOM	299	CG	GLU	A	53	32.189	22.603	8.081	1.00	35.02	C
ATOM	300	CD	GLU	A	53	31.785	21.185	8.507	1.00	37.87	C
ATOM	301	OE1	GLU	A	53	30.614	20.929	8.887	1.00	39.16	O
ATOM	302	OE2	GLU	A	53	32.674	20.313	8.463	1.00	40.43	O
ATOM	303	N	GLU	A	54	30.811	26.479	7.652	1.00	29.84	N
ATOM	304	CA	GLU	A	54	31.413	27.796	7.757	1.00	30.22	C
ATOM	305	C	GLU	A	54	30.975	28.486	9.045	1.00	29.05	C
ATOM	306	O	GLU	A	54	31.780	29.102	9.719	1.00	28.03	O
ATOM	307	CB	GLU	A	54	31.026	28.680	6.573	1.00	30.53	C
ATOM	308	CG	GLU	A	54	31.635	28.276	5.243	1.00	34.66	C
ATOM	309	CD	GLU	A	54	30.993	29.002	4.058	1.00	38.49	C
ATOM	310	OE1	GLU	A	54	30.651	30.208	4.214	1.00	42.20	O
ATOM	311	OE2	GLU	A	54	30.829	28.368	2.975	1.00	41.54	O
ATOM	312	N	LEU	A	55	29.696	28.366	9.387	1.00	28.24	N
ATOM	313	CA	LEU	A	55	29.177	29.029	10.571	1.00	27.28	C
ATOM	314	C	LEU	A	55	29.774	28.448	11.827	1.00	26.43	C
ATOM	315	O	LEU	A	55	30.212	29.199	12.695	1.00	25.20	O
ATOM	316	CB	LEU	A	55	27.655	28.975	10.597	1.00	27.25	C

ATOM	317	CG	LEU	A	55	27.027	29.829	9.502	1.00	27.43
ATOM	318	CD1	LEU	A	55	25.568	29.452	9.322	1.00	28.83
ATOM	319	CD2	LEU	A	55	27.173	31.318	9.821	1.00	26.98
ATOM	320	N	ILE	A	56	29.829	27.116	11.915	1.00	25.75
ATOM	321	CA	ILE	A	56	30.382	26.473	13.107	1.00	25.50
ATOM	322	C	ILE	A	56	31.874	26.815	13.254	1.00	26.26
ATOM	323	O	ILE	A	56	32.346	27.134	14.349	1.00	25.55
ATOM	324	CB	ILE	A	56	30.192	24.955	13.059	1.00	25.63
ATOM	325	CG1	ILE	A	56	28.698	24.574	13.034	1.00	24.00
ATOM	326	CG2	ILE	A	56	30.848	24.321	14.263	1.00	25.48
ATOM	327	CD1	ILE	A	56	28.439	23.170	12.531	1.00	24.10
ATOM	328	N	GLU	A	57	32.597	26.781	12.135	1.00	27.04
ATOM	329	CA	GLU	A	57	34.023	27.082	12.107	1.00	28.01
ATOM	330	C	GLU	A	57	34.232	28.491	12.625	1.00	28.49
ATOM	331	O	GLU	A	57	35.183	28.770	13.344	1.00	28.92
ATOM	332	CB	GLU	A	57	34.561	26.977	10.676	1.00	28.61
ATOM	333	CG	GLU	A	57	36.053	27.224	10.509	1.00	30.77
ATOM	334	CD	GLU	A	57	36.902	26.322	11.394	1.00	35.15
ATOM	335	OE1	GLU	A	57	36.556	25.127	11.591	1.00	37.11
ATOM	336	OE2	GLU	A	57	37.924	26.818	11.909	1.00	40.88
ATOM	337	N	ASN	A	58	33.321	29.378	12.270	1.00	28.34
ATOM	338	CA	ASN	A	58	33.424	30.758	12.701	1.00	28.81
ATOM	339	C	ASN	A	58	32.770	31.079	14.025	1.00	27.23
ATOM	340	O	ASN	A	58	32.630	32.233	14.374	1.00	26.07
ATOM	341	CB	ASN	A	58	32.792	31.641	11.656	1.00	29.66
ATOM	342	CG	ASN	A	58	33.789	32.386	10.913	1.00	33.61
ATOM	343	OD1	ASN	A	58	34.280	31.901	9.893	1.00	38.81
ATOM	344	ND2	ASN	A	58	34.160	33.582	11.429	1.00	36.72
ATOM	345	N	GLU	A	59	32.343	30.053	14.735	1.00	26.39
ATOM	346	CA	GLU	A	59	31.712	30.241	16.030	1.00	26.36
ATOM	347	C	GLU	A	59	30.495	31.144	15.926	1.00	25.39
ATOM	348	O	GLU	A	59	30.325	32.100	16.668	1.00	24.14
ATOM	349	CB	GLU	A	59	32.753	30.697	17.059	1.00	26.80
ATOM	350	CG	GLU	A	59	33.717	29.537	17.316	1.00	29.10
ATOM	351	CD	GLU	A	59	34.722	29.791	18.407	1.00	32.85
ATOM	352	OE1	GLU	A	59	35.790	30.330	18.080	1.00	37.47
ATOM	353	OE2	GLU	A	59	34.466	29.432	19.572	1.00	34.14
ATOM	354	N	GLU	A	60	29.641	30.799	14.970	1.00	24.88
ATOM	355	CA	GLU	A	60	28.366	31.465	14.801	1.00	25.20
ATOM	356	C	GLU	A	60	27.262	30.414	14.822	1.00	24.07
ATOM	357	O	GLU	A	60	27.420	29.320	14.293	1.00	22.49
ATOM	358	CB	GLU	A	60	28.343	32.249	13.505	1.00	25.69
ATOM	359	CG	GLU	A	60	29.354	33.363	13.536	1.00	30.40
ATOM	360	CD	GLU	A	60	28.962	34.516	12.657	1.00	37.91
ATOM	361	OE1	GLU	A	60	29.009	34.377	11.414	1.00	38.24
ATOM	362	OE2	GLU	A	60	28.595	35.566	13.236	1.00	46.50
ATOM	363	N	PRO	A	61	26.124	30.779	15.386	1.00	22.90
ATOM	364	CA	PRO	A	61	25.045	29.816	15.571	1.00	22.19
ATOM	365	C	PRO	A	61	24.448	29.371	14.267	1.00	21.47
ATOM	366	O	PRO	A	61	24.419	30.104	13.266	1.00	20.48
ATOM	367	CB	PRO	A	61	23.996	30.583	16.368	1.00	22.29
ATOM	368	CG	PRO	A	61	24.315	32.053	16.148	1.00	22.67
ATOM	369	CD	PRO	A	61	25.764	32.140	15.798	1.00	22.92
ATOM	370	N	VAL	A	62	23.957	28.142	14.283	1.00	20.87
ATOM	371	CA	VAL	A	62	23.248	27.591	13.146	1.00	20.94
ATOM	372	C	VAL	A	62	22.353	26.475	13.655	1.00	21.15
ATOM	373	O	VAL	A	62	22.714	25.761	14.605	1.00	21.21
ATOM	374	CB	VAL	A	62	24.214	27.052	12.073	1.00	21.30
ATOM	375	CG1	VAL	A	62	25.061	25.912	12.608	1.00	22.10
ATOM	376	CG2	VAL	A	62	23.440	26.587	10.825	1.00	21.19
ATOM	377	N	VAL	A	63	21.158	26.366	13.084	1.00	21.48

C
C
C
N
C
C
C
O
C
C
C
N
C
C
C
O
O
N
C
C
C
O
C
C
C
O
N
C
C
C
O
C
C
C
C
C
C
N
C
C
C
O
C
C
C
N

ATOM	378	CA	VAL	A	63	20.302	25.231	13.374	1.00	21.87	C
ATOM	379	C	VAL	A	63	20.526	24.212	12.270	1.00	22.00	C
ATOM	380	O	VAL	A	63	20.366	24.528	11.109	1.00	22.54	O
ATOM	381	CB	VAL	A	63	18.793	25.574	13.421	1.00	22.02	C
ATOM	382	CG1	VAL	A	63	17.955	24.277	13.612	1.00	21.90	C
ATOM	383	CG2	VAL	A	63	18.494	26.522	14.546	1.00	21.22	C
ATOM	384	N	LEU	A	64	20.911	23.002	12.643	1.00	22.48	N
ATOM	385	CA	LEU	A	64	21.067	21.902	11.710	1.00	23.08	C
ATOM	386	C	LEU	A	64	19.826	21.014	11.775	1.00	22.19	C
ATOM	387	O	LEU	A	64	19.423	20.558	12.843	1.00	21.85	O
ATOM	388	CB	LEU	A	64	22.327	21.107	12.032	1.00	24.00	C
ATOM	389	CG	LEU	A	64	23.614	21.930	11.984	1.00	27.02	C
ATOM	390	CD1	LEU	A	64	24.791	21.080	12.303	1.00	31.61	C
ATOM	391	CD2	LEU	A	64	23.827	22.540	10.640	1.00	29.86	C
ATOM	392	N	THR	A	65	19.222	20.757	10.628	1.00	21.50	N
ATOM	393	CA	THR	A	65	17.943	20.056	10.613	1.00	21.95	C
ATOM	394	C	THR	A	65	18.022	18.561	10.389	1.00	21.35	C
ATOM	395	O	THR	A	65	17.028	17.870	10.581	1.00	21.37	O
ATOM	396	CB	THR	A	65	17.062	20.592	9.478	1.00	22.08	C
ATOM	397	OG1	THR	A	65	17.725	20.351	8.230	1.00	22.38	O
ATOM	398	CG2	THR	A	65	16.919	22.099	9.553	1.00	24.00	C
ATOM	399	N	ASP	A	66	19.168	18.062	9.961	1.00	21.50	N
ATOM	400	CA	ASP	A	66	19.259	16.650	9.604	1.00	21.86	C
ATOM	401	C	ASP	A	66	20.513	15.899	10.069	1.00	21.61	C
ATOM	402	O	ASP	A	66	21.070	15.088	9.316	1.00	21.82	O
ATOM	403	CB	ASP	A	66	19.152	16.553	8.084	1.00	22.20	C
ATOM	404	CG	ASP	A	66	20.199	17.355	7.390	1.00	21.49	C
ATOM	405	OD1	ASP	A	66	21.065	17.936	8.070	1.00	21.14	O
ATOM	406	OD2	ASP	A	66	20.240	17.477	6.159	1.00	25.40	O
ATOM	407	N	THR	A	67	20.967	16.158	11.287	1.00	21.05	N
ATOM	408	CA	THR	A	67	22.154	15.491	11.768	1.00	20.85	C
ATOM	409	C	THR	A	67	21.858	14.084	12.231	1.00	20.12	C
ATOM	410	O	THR	A	67	22.757	13.281	12.283	1.00	20.45	O
ATOM	411	CB	THR	A	67	22.747	16.207	12.977	1.00	20.77	C
ATOM	412	OG1	THR	A	67	21.782	16.210	14.039	1.00	20.61	O
ATOM	413	CG2	THR	A	67	23.054	17.671	12.682	1.00	22.44	C
ATOM	414	N	ASN	A	68	20.614	13.815	12.618	1.00	19.51	N
ATOM	415	CA	ASN	A	68	20.269	12.555	13.267	1.00	19.09	C
ATOM	416	C	ASN	A	68	21.116	12.349	14.517	1.00	18.40	C
ATOM	417	O	ASN	A	68	21.383	11.212	14.928	1.00	17.67	O
ATOM	418	CB	ASN	A	68	20.455	11.362	12.325	1.00	19.63	C
ATOM	419	CG	ASN	A	68	19.453	11.344	11.209	1.00	19.93	C
ATOM	420	OD1	ASN	A	68	18.253	11.236	11.442	1.00	19.76	O
ATOM	421	ND2	ASN	A	68	19.941	11.423	9.987	1.00	19.96	N
ATOM	422	N	LEU	A	69	21.532	13.448	15.134	1.00	18.28	N
ATOM	423	CA	LEU	A	69	22.378	13.382	16.326	1.00	17.98	C
ATOM	424	C	LEU	A	69	21.773	12.542	17.447	1.00	17.59	C
ATOM	425	O	LEU	A	69	22.478	11.725	18.039	1.00	17.61	O
ATOM	426	CB	LEU	A	69	22.693	14.772	16.844	1.00	18.38	C
ATOM	427	CG	LEU	A	69	23.636	14.859	18.035	1.00	18.31	C
ATOM	428	CD1	LEU	A	69	24.936	14.144	17.744	1.00	20.68	C
ATOM	429	CD2	LEU	A	69	23.907	16.299	18.399	1.00	19.25	C
ATOM	430	N	VAL	A	70	20.489	12.741	17.740	1.00	17.27	N
ATOM	431	CA	VAL	A	70	19.811	11.989	18.794	1.00	17.72	C
ATOM	432	C	VAL	A	70	18.588	11.253	18.257	1.00	18.11	C
ATOM	433	O	VAL	A	70	17.557	11.148	18.927	1.00	17.23	O
ATOM	434	CB	VAL	A	70	19.395	12.866	19.999	1.00	17.54	C
ATOM	435	CG1	VAL	A	70	20.624	13.402	20.704	1.00	19.07	C
ATOM	436	CG2	VAL	A	70	18.451	13.994	19.599	1.00	17.50	C
ATOM	437	N	TYR	A	71	18.715	10.764	17.028	1.00	19.22	N
ATOM	438	CA	TYR	A	71	17.615	10.071	16.367	1.00	19.79	C

ATOM	500	N	LEU	A	78	7.790	10.705	28.775	1.00	25.96	N
ATOM	501	CA	LEU	A	78	7.893	10.441	30.201	1.00	26.27	C
ATOM	502	C	LEU	A	78	7.716	8.965	30.539	1.00	26.14	C
ATOM	503	O	LEU	A	78	8.446	8.438	31.374	1.00	25.51	O
ATOM	504	CB	LEU	A	78	6.905	11.301	30.979	1.00	26.32	C
ATOM	505	CG	LEU	A	78	7.155	12.802	30.860	1.00	27.35	C
ATOM	506	CD1	LEU	A	78	6.098	13.579	31.623	1.00	28.05	C
ATOM	507	CD2	LEU	A	78	8.536	13.165	31.379	1.00	28.39	C
ATOM	508	N	GLU	A	79	6.775	8.297	29.874	1.00	26.56	N
ATOM	509	CA	GLU	A	79	6.526	6.876	30.123	1.00	27.00	C
ATOM	510	C	GLU	A	79	7.754	6.029	29.780	1.00	26.63	C
ATOM	511	O	GLU	A	79	8.220	5.231	30.583	1.00	25.94	O
ATOM	512	CB	GLU	A	79	5.302	6.371	29.333	1.00	27.39	C
ATOM	513	CG	GLU	A	79	4.963	4.919	29.657	1.00	29.75	C
ATOM	514	CD	GLU	A	79	3.803	4.347	28.854	1.00	33.15	C
ATOM	515	OE1	GLU	A	79	3.361	4.976	27.863	1.00	35.39	O
ATOM	516	OE2	GLU	A	79	3.338	3.245	29.224	1.00	35.44	O
ATOM	517	N	TYR	A	80	8.279	6.205	28.575	1.00	26.53	N
ATOM	518	CA	TYR	A	80	9.466	5.465	28.169	1.00	26.29	C
ATOM	519	C	TYR	A	80	10.669	5.763	29.072	1.00	25.83	C
ATOM	520	O	TYR	A	80	11.418	4.863	29.435	1.00	25.75	O
ATOM	521	CB	TYR	A	80	9.812	5.802	26.728	1.00	26.53	C
ATOM	522	CG	TYR	A	80	11.047	5.106	26.210	1.00	26.94	C
ATOM	523	CD1	TYR	A	80	10.971	3.836	25.637	1.00	26.53	C
ATOM	524	CD2	TYR	A	80	12.287	5.721	26.291	1.00	25.32	C
ATOM	525	CE1	TYR	A	80	12.113	3.208	25.142	1.00	26.28	C
ATOM	526	CE2	TYR	A	80	13.416	5.109	25.823	1.00	25.73	C
ATOM	527	CZ	TYR	A	80	13.331	3.861	25.240	1.00	25.94	C
ATOM	528	OH	TYR	A	80	14.478	3.294	24.758	1.00	25.55	O
ATOM	529	N	LEU	A	81	10.859	7.016	29.448	1.00	25.62	N
ATOM	530	CA	LEU	A	81	12.000	7.350	30.309	1.00	25.87	C
ATOM	531	C	LEU	A	81	11.821	6.794	31.733	1.00	26.06	C
ATOM	532	O	LEU	A	81	12.763	6.263	32.327	1.00	26.00	O
ATOM	533	CB	LEU	A	81	12.250	8.863	30.340	1.00	25.81	C
ATOM	534	CG	LEU	A	81	12.748	9.501	29.030	1.00	25.65	C
ATOM	535	CD1	LEU	A	81	12.828	11.010	29.174	1.00	26.40	C
ATOM	536	CD2	LEU	A	81	14.102	8.959	28.592	1.00	25.79	C
ATOM	537	N	GLN	A	82	10.617	6.913	32.274	1.00	26.27	N
ATOM	538	CA	GLN	A	82	10.338	6.388	33.602	1.00	26.77	C
ATOM	539	C	GLN	A	82	10.640	4.897	33.623	1.00	26.51	C
ATOM	540	O	GLN	A	82	11.232	4.384	34.552	1.00	26.60	O
ATOM	541	CB	GLN	A	82	8.877	6.611	33.974	1.00	26.90	C
ATOM	542	CG	GLN	A	82	8.466	5.856	35.236	1.00	28.41	C
ATOM	543	CD	GLN	A	82	7.315	6.501	35.987	1.00	30.63	C
ATOM	544	OE1	GLN	A	82	6.755	7.501	35.551	1.00	32.13	O
ATOM	545	NE2	GLN	A	82	6.944	5.907	37.116	1.00	34.12	N
ATOM	546	N	GLU	A	83	10.252	4.216	32.561	1.00	26.68	N
ATOM	547	CA	GLU	A	83	10.429	2.773	32.456	1.00	26.90	C
ATOM	548	C	GLU	A	83	11.893	2.359	32.295	1.00	26.42	C
ATOM	549	O	GLU	A	83	12.285	1.294	32.764	1.00	25.50	O
ATOM	550	CB	GLU	A	83	9.604	2.257	31.273	1.00	26.84	C
ATOM	551	CG	GLU	A	83	9.607	0.753	31.080	1.00	29.20	C
ATOM	552	CD	GLU	A	83	8.902	0.014	32.205	1.00	31.92	C
ATOM	553	OE1	GLU	A	83	8.258	0.666	33.060	1.00	34.06	O
ATOM	554	OE2	GLU	A	83	9.008	-1.221	32.243	1.00	33.78	O
ATOM	555	N	ASN	A	84	12.711	3.223	31.700	1.00	25.79	N
ATOM	556	CA	ASN	A	84	14.063	2.824	31.312	1.00	25.87	C
ATOM	557	C	ASN	A	84	15.273	3.607	31.810	1.00	26.08	C
ATOM	558	O	ASN	A	84	16.392	3.138	31.628	1.00	26.05	O
ATOM	559	CB	ASN	A	84	14.137	2.844	29.784	1.00	25.74	C
ATOM	560	CG	ASN	A	84	13.262	1.799	29.152	1.00	25.67	C

ATOM	561	OD1	ASN	A	84	13.402	0.608	29.439	1.00	24.26
ATOM	562	ND2	ASN	A	84	12.354	2.231	28.288	1.00	22.66
ATOM	563	N	ILE	A	85	15.087	4.773	32.422	1.00	26.02
ATOM	564	CA	ILE	A	85	16.240	5.599	32.763	1.00	26.74
ATOM	565	C	ILE	A	85	16.930	5.307	34.099	1.00	26.89
ATOM	566	O	ILE	A	85	17.820	6.054	34.519	1.00	26.93
ATOM	567	CB	ILE	A	85	15.846	7.076	32.688	1.00	26.70
ATOM	568	CG1	ILE	A	85	17.049	7.923	32.270	1.00	27.99
ATOM	569	CG2	ILE	A	85	15.232	7.535	33.988	1.00	26.50
ATOM	570	CD1	ILE	A	85	16.696	9.389	32.000	1.00	28.77
ATOM	571	N	GLY	A	86	16.503	4.253	34.773	1.00	27.27
ATOM	572	CA	GLY	A	86	17.130	3.834	36.018	1.00	27.51
ATOM	573	C	GLY	A	86	16.573	4.453	37.289	1.00	27.73
ATOM	574	O	GLY	A	86	15.563	5.163	37.268	1.00	27.80
ATOM	575	N	ASN	A	87	17.273	4.186	38.392	1.00	27.70
ATOM	576	CA	ASN	A	87	16.864	4.603	39.726	1.00	27.53
ATOM	577	C	ASN	A	87	17.849	5.602	40.320	1.00	27.33
ATOM	578	O	ASN	A	87	17.932	5.762	41.539	1.00	26.97
ATOM	579	CB	ASN	A	87	16.726	3.367	40.644	1.00	27.60
ATOM	580	N	GLY	A	88	18.601	6.273	39.455	1.00	27.41
ATOM	581	CA	GLY	A	88	19.528	7.307	39.888	1.00	27.29
ATOM	582	C	GLY	A	88	18.762	8.549	40.288	1.00	27.18
ATOM	583	O	GLY	A	88	17.570	8.654	40.001	1.00	27.43
ATOM	584	N	ASP	A	89	19.437	9.489	40.945	1.00	27.04
ATOM	585	CA	ASP	A	89	18.808	10.739	41.366	1.00	26.88
ATOM	586	C	ASP	A	89	18.761	11.755	40.227	1.00	26.69
ATOM	587	O	ASP	A	89	19.697	11.848	39.446	1.00	26.91
ATOM	588	CB	ASP	A	89	19.596	11.372	42.504	1.00	26.95
ATOM	589	CG	ASP	A	89	19.375	10.688	43.834	1.00	27.28
ATOM	590	OD1	ASP	A	89	18.499	9.811	43.939	1.00	29.31
ATOM	591	OD2	ASP	A	89	20.040	10.983	44.839	1.00	27.32
ATOM	592	N	PHE	A	90	17.687	12.531	40.152	1.00	26.24
ATOM	593	CA	PHE	A	90	17.598	13.594	39.165	1.00	26.10
ATOM	594	C	PHE	A	90	17.407	14.931	39.866	1.00	26.23
ATOM	595	O	PHE	A	90	16.629	15.037	40.811	1.00	26.48
ATOM	596	CB	PHE	A	90	16.460	13.322	38.184	1.00	25.91
ATOM	597	CG	PHE	A	90	16.747	12.201	37.241	1.00	25.26
ATOM	598	CD1	PHE	A	90	16.601	10.886	37.640	1.00	25.32
ATOM	599	CD2	PHE	A	90	17.190	12.459	35.958	1.00	26.02
ATOM	600	CE1	PHE	A	90	16.872	9.856	36.777	1.00	24.09
ATOM	601	CE2	PHE	A	90	17.460	11.427	35.091	1.00	25.07
ATOM	602	CZ	PHE	A	90	17.308	10.117	35.512	1.00	24.67
ATOM	603	N	SER	A	91	18.149	15.941	39.430	1.00	26.33
ATOM	604	CA	SER	A	91	17.985	17.280	39.973	1.00	26.48
ATOM	605	C	SER	A	91	16.777	17.923	39.310	1.00	26.71
ATOM	606	O	SER	A	91	16.696	18.029	38.077	1.00	26.09
ATOM	607	CB	SER	A	91	19.226	18.152	39.763	1.00	26.25
ATOM	608	OG	SER	A	91	20.364	17.543	40.326	1.00	26.61
ATOM	609	N	VAL	A	92	15.835	18.327	40.150	1.00	27.06
ATOM	610	CA	VAL	A	92	14.629	18.971	39.695	1.00	27.44
ATOM	611	C	VAL	A	92	14.468	20.295	40.405	1.00	27.57
ATOM	612	O	VAL	A	92	14.475	20.376	41.642	1.00	26.69
ATOM	613	CB	VAL	A	92	13.407	18.132	40.004	1.00	27.60
ATOM	614	CG1	VAL	A	92	12.164	18.800	39.432	1.00	28.15
ATOM	615	CG2	VAL	A	92	13.584	16.731	39.457	1.00	27.84
ATOM	616	N	TYR	A	93	14.312	21.334	39.598	1.00	27.77
ATOM	617	CA	TYR	A	93	14.120	22.661	40.108	1.00	27.91
ATOM	618	C	TYR	A	93	12.654	22.940	40.172	1.00	28.16
ATOM	619	O	TYR	A	93	11.894	22.525	39.303	1.00	28.02
ATOM	620	CB	TYR	A	93	14.810	23.673	39.216	1.00	27.94
ATOM	621	CG	TYR	A	93	16.291	23.593	39.368	1.00	29.53

ATOM	622	CD1	TYR	A	93	16.929	24.258	40.395	1.00	29.48	C
ATOM	623	CD2	TYR	A	93	17.046	22.815	38.522	1.00	30.95	C
ATOM	624	CE1	TYR	A	93	18.266	24.169	40.553	1.00	29.93	C
ATOM	625	CE2	TYR	A	93	18.386	22.726	38.679	1.00	31.90	C
ATOM	626	CZ	TYR	A	93	18.985	23.402	39.704	1.00	31.56	C
ATOM	627	OH	TYR	A	93	20.327	23.306	39.878	1.00	35.47	O
ATOM	628	N	SER	A	94	12.282	23.669	41.208	1.00	28.70	N
ATOM	629	CA	SER	A	94	10.913	24.037	41.459	1.00	29.47	C
ATOM	630	C	SER	A	94	10.856	25.547	41.554	1.00	29.69	C
ATOM	631	O	SER	A	94	11.705	26.167	42.187	1.00	29.50	O
ATOM	632	CB	SER	A	94	10.456	23.411	42.774	1.00	29.32	C
ATOM	633	OG	SER	A	94	9.093	23.701	43.009	1.00	30.85	O
ATOM	634	N	ALA	A	95	9.858	26.146	40.928	1.00	30.39	N
ATOM	635	CA	ALA	A	95	9.760	27.591	40.929	1.00	31.04	C
ATOM	636	C	ALA	A	95	8.330	28.064	40.932	1.00	31.76	C
ATOM	637	O	ALA	A	95	7.425	27.386	40.452	1.00	31.87	O
ATOM	638	CB	ALA	A	95	10.480	28.165	39.716	1.00	31.12	C
ATOM	639	N	SER	A	96	8.147	29.258	41.467	1.00	32.54	N
ATOM	640	CA	SER	A	96	6.848	29.891	41.489	1.00	33.42	C
ATOM	641	C	SER	A	96	6.708	30.869	40.326	1.00	33.15	C
ATOM	642	O	SER	A	96	5.662	31.487	40.163	1.00	34.35	O
ATOM	643	CB	SER	A	96	6.673	30.661	42.787	1.00	33.88	C
ATOM	644	OG	SER	A	96	5.380	31.217	42.815	1.00	35.79	O
ATOM	645	N	THR	A	97	7.775	31.031	39.554	1.00	32.09	N
ATOM	646	CA	THR	A	97	7.797	31.920	38.397	1.00	31.44	C
ATOM	647	C	THR	A	97	8.185	31.095	37.191	1.00	30.26	C
ATOM	648	O	THR	A	97	8.693	30.003	37.348	1.00	29.72	O
ATOM	649	CB	THR	A	97	8.857	33.031	38.582	1.00	31.64	C
ATOM	650	OG1	THR	A	97	9.147	33.656	37.323	1.00	32.17	O
ATOM	651	CG2	THR	A	97	10.227	32.455	38.978	1.00	31.73	C
ATOM	652	N	HIS	A	98	7.974	31.623	35.996	1.00	29.35	N
ATOM	653	CA	HIS	A	98	8.379	30.915	34.780	1.00	28.83	C
ATOM	654	C	HIS	A	98	9.899	30.969	34.589	1.00	28.37	C
ATOM	655	O	HIS	A	98	10.467	30.189	33.824	1.00	27.53	O
ATOM	656	CB	HIS	A	98	7.671	31.490	33.548	1.00	28.76	C
ATOM	657	CG	HIS	A	98	7.936	32.942	33.309	1.00	28.53	C
ATOM	658	ND1	HIS	A	98	7.271	33.944	33.984	1.00	29.18	N
ATOM	659	CD2	HIS	A	98	8.793	33.566	32.463	1.00	29.68	C
ATOM	660	CE1	HIS	A	98	7.708	35.120	33.568	1.00	29.42	C
ATOM	661	NE2	HIS	A	98	8.638	34.919	32.650	1.00	29.10	N
ATOM	662	N	LYS	A	99	10.562	31.874	35.299	1.00	27.85	N
ATOM	663	CA	LYS	A	99	11.997	32.041	35.124	1.00	28.24	C
ATOM	664	C	LYS	A	99	12.878	31.143	35.997	1.00	27.69	C
ATOM	665	O	LYS	A	99	12.892	31.278	37.216	1.00	29.10	O
ATOM	666	CB	LYS	A	99	12.366	33.498	35.349	1.00	28.47	C
ATOM	667	CG	LYS	A	99	11.836	34.410	34.255	1.00	30.42	C
ATOM	668	CD	LYS	A	99	12.431	35.791	34.347	1.00	32.66	C
ATOM	669	CE	LYS	A	99	11.757	36.624	35.421	1.00	34.45	C
ATOM	670	NZ	LYS	A	99	10.569	37.347	34.884	1.00	35.39	N
ATOM	671	N	PHE	A	100	13.621	30.237	35.376	1.00	26.40	N
ATOM	672	CA	PHE	A	100	14.551	29.397	36.127	1.00	25.95	C
ATOM	673	C	PHE	A	100	15.978	29.920	35.995	1.00	25.87	C
ATOM	674	O	PHE	A	100	16.809	29.386	35.247	1.00	25.86	O
ATOM	675	CB	PHE	A	100	14.469	27.945	35.686	1.00	25.48	C
ATOM	676	CG	PHE	A	100	13.211	27.267	36.109	1.00	25.06	C
ATOM	677	CD1	PHE	A	100	12.047	27.436	35.388	1.00	24.80	C
ATOM	678	CD2	PHE	A	100	13.191	26.472	37.238	1.00	25.18	C
ATOM	679	CE1	PHE	A	100	10.886	26.805	35.772	1.00	25.96	C
ATOM	680	CE2	PHE	A	100	12.030	25.835	37.631	1.00	25.79	C
ATOM	681	CZ	PHE	A	100	10.878	26.004	36.898	1.00	26.05	C
ATOM	682	N	LEU	A	101	16.237	30.997	36.709	1.00	25.77	N

ATOM	683	CA	LEU	A	101	17.549	31.589	36.747	1.00	25.85	C
ATOM	684	C	LEU	A	101	18.527	30.605	37.381	1.00	25.87	C
ATOM	685	O	LEU	A	101	18.319	30.136	38.503	1.00	24.99	O
ATOM	686	CB	LEU	A	101	17.488	32.876	37.559	1.00	25.84	C
ATOM	687	CG	LEU	A	101	18.795	33.656	37.708	1.00	26.07	C
ATOM	688	CD1	LEU	A	101	19.244	34.200	36.377	1.00	25.11	C
ATOM	689	CD2	LEU	A	101	18.609	34.800	38.710	1.00	27.75	C
ATOM	690	N	TYR	A	102	19.582	30.274	36.644	1.00	26.56	N
ATOM	691	CA	TYR	A	102	20.611	29.380	37.159	1.00	26.99	C
ATOM	692	C	TYR	A	102	21.430	30.072	38.260	1.00	27.10	C
ATOM	693	O	TYR	A	102	21.711	31.264	38.174	1.00	27.00	O
ATOM	694	CB	TYR	A	102	21.562	28.939	36.053	1.00	27.26	C
ATOM	695	CG	TYR	A	102	22.667	28.088	36.620	1.00	28.93	C
ATOM	696	CD1	TYR	A	102	22.457	26.750	36.914	1.00	29.52	C
ATOM	697	CD2	TYR	A	102	23.906	28.636	36.909	1.00	31.11	C
ATOM	698	CE1	TYR	A	102	23.463	25.979	37.480	1.00	32.23	C
ATOM	699	CE2	TYR	A	102	24.913	27.872	37.470	1.00	32.52	C
ATOM	700	CZ	TYR	A	102	24.693	26.549	37.750	1.00	33.88	C
ATOM	701	OH	TYR	A	102	25.727	25.794	38.314	1.00	38.68	O
ATOM	702	N	TYR	A	103	21.793	29.322	39.296	1.00	26.83	N
ATOM	703	CA	TYR	A	103	22.667	29.840	40.343	1.00	27.22	C
ATOM	704	C	TYR	A	103	23.613	28.750	40.834	1.00	26.77	C
ATOM	705	O	TYR	A	103	23.287	27.556	40.853	1.00	25.60	O
ATOM	706	CB	TYR	A	103	21.880	30.440	41.517	1.00	27.38	C
ATOM	707	CG	TYR	A	103	20.909	29.493	42.154	1.00	29.11	C
ATOM	708	CD1	TYR	A	103	19.639	29.340	41.639	1.00	30.58	C
ATOM	709	CD2	TYR	A	103	21.257	28.759	43.279	1.00	32.13	C
ATOM	710	CE1	TYR	A	103	18.734	28.469	42.212	1.00	32.78	C
ATOM	711	CE2	TYR	A	103	20.357	27.887	43.877	1.00	33.15	C
ATOM	712	CZ	TYR	A	103	19.096	27.745	43.329	1.00	34.41	C
ATOM	713	OH	TYR	A	103	18.187	26.888	43.892	1.00	37.44	O
ATOM	714	N	ASP	A	104	24.798	29.182	41.225	1.00	26.36	N
ATOM	715	CA	ASP	A	104	25.832	28.273	41.673	1.00	26.28	C
ATOM	716	C	ASP	A	104	25.802	28.239	43.184	1.00	26.21	C
ATOM	717	O	ASP	A	104	26.163	29.212	43.845	1.00	25.45	O
ATOM	718	CB	ASP	A	104	27.174	28.765	41.154	1.00	25.92	C
ATOM	719	CG	ASP	A	104	28.330	27.889	41.575	1.00	27.39	C
ATOM	720	OD1	ASP	A	104	28.161	26.973	42.437	1.00	27.13	O
ATOM	721	OD2	ASP	A	104	29.459	28.068	41.075	1.00	27.68	O
ATOM	722	N	GLU	A	105	25.358	27.114	43.724	1.00	26.46	N
ATOM	723	CA	GLU	A	105	25.218	26.962	45.162	1.00	27.02	C
ATOM	724	C	GLU	A	105	26.540	27.199	45.916	1.00	26.91	C
ATOM	725	O	GLU	A	105	26.523	27.718	47.026	1.00	26.51	O
ATOM	726	CB	GLU	A	105	24.602	25.587	45.481	1.00	27.45	C
ATOM	727	CG	GLU	A	105	23.088	25.556	45.254	1.00	29.46	C
ATOM	728	CD	GLU	A	105	22.527	24.158	45.025	1.00	32.85	C
ATOM	729	OE1	GLU	A	105	22.908	23.234	45.765	1.00	32.90	O
ATOM	730	OE2	GLU	A	105	21.703	23.980	44.085	1.00	35.16	O
ATOM	731	N	LYS	A	106	27.680	26.874	45.305	1.00	27.12	N
ATOM	732	CA	LYS	A	106	28.979	27.041	45.983	1.00	27.54	C
ATOM	733	C	LYS	A	106	29.331	28.493	46.258	1.00	27.70	C
ATOM	734	O	LYS	A	106	30.164	28.769	47.099	1.00	27.61	O
ATOM	735	CB	LYS	A	106	30.127	26.441	45.163	1.00	27.58	C
ATOM	736	CG	LYS	A	106	30.038	24.928	44.918	1.00	28.61	C
ATOM	737	N	LYS	A	107	28.713	29.420	45.537	1.00	27.99	N
ATOM	738	CA	LYS	A	107	29.020	30.826	45.694	1.00	28.30	C
ATOM	739	C	LYS	A	107	28.034	31.513	46.631	1.00	29.21	C
ATOM	740	O	LYS	A	107	28.175	32.701	46.917	1.00	29.08	O
ATOM	741	CB	LYS	A	107	29.040	31.524	44.324	1.00	28.30	C
ATOM	742	CG	LYS	A	107	30.344	31.291	43.516	1.00	28.07	C
ATOM	743	CD	LYS	A	107	30.242	31.737	42.034	1.00	27.29	C

ATOM	805	CZ	PHE	A	114	20.066	33.866	43.172	1.00	40.52
ATOM	806	N	LYS	A	115	18.669	35.457	47.461	1.00	39.11
ATOM	807	CA	LYS	A	115	17.596	34.529	47.787	1.00	39.02
ATOM	808	C	LYS	A	115	17.042	33.938	46.497	1.00	38.71
ATOM	809	O	LYS	A	115	16.285	34.590	45.787	1.00	39.00
ATOM	810	CB	LYS	A	115	16.498	35.251	48.578	1.00	39.14
ATOM	811	N	PRO	A	116	17.389	32.691	46.213	1.00	38.17
ATOM	812	CA	PRO	A	116	17.027	32.060	44.938	1.00	37.95
ATOM	813	C	PRO	A	116	15.532	31.889	44.783	1.00	37.38
ATOM	814	O	PRO	A	116	14.865	31.506	45.733	1.00	37.36
ATOM	815	CB	PRO	A	116	17.684	30.679	45.004	1.00	38.15
ATOM	816	CG	PRO	A	116	18.416	30.604	46.297	1.00	38.40
ATOM	817	CD	PRO	A	116	18.079	31.775	47.125	1.00	38.19
ATOM	818	N	ARG	A	117	15.027	32.161	43.590	1.00	36.84
ATOM	819	CA	ARG	A	117	13.611	32.028	43.293	1.00	36.61
ATOM	820	C	ARG	A	117	13.259	30.581	42.984	1.00	36.38
ATOM	821	O	ARG	A	117	12.084	30.230	42.913	1.00	36.44
ATOM	822	CB	ARG	A	117	13.235	32.913	42.122	1.00	36.62
ATOM	823	N	SER	A	118	14.271	29.745	42.777	1.00	35.79
ATOM	824	CA	SER	A	118	14.025	28.339	42.538	1.00	35.51
ATOM	825	C	SER	A	118	14.844	27.491	43.512	1.00	35.35
ATOM	826	O	SER	A	118	15.888	27.923	43.991	1.00	35.16
ATOM	827	CB	SER	A	118	14.330	27.976	41.086	1.00	35.26
ATOM	828	OG	SER	A	118	15.689	28.164	40.795	1.00	35.14
ATOM	829	N	ASN	A	119	14.329	26.306	43.824	1.00	35.11
ATOM	830	CA	ASN	A	119	14.997	25.373	44.715	1.00	35.21
ATOM	831	C	ASN	A	119	15.233	24.049	44.022	1.00	34.26
ATOM	832	O	ASN	A	119	14.401	23.595	43.244	1.00	34.11
ATOM	833	CB	ASN	A	119	14.145	25.110	45.957	1.00	35.95
ATOM	834	CG	ASN	A	119	13.636	26.382	46.592	1.00	38.47
ATOM	835	OD1	ASN	A	119	14.417	27.245	47.025	1.00	41.76
ATOM	836	ND2	ASN	A	119	12.314	26.511	46.657	1.00	42.21
ATOM	837	N	ARG	A	120	16.368	23.434	44.326	1.00	33.75
ATOM	838	CA	ARG	A	120	16.750	22.149	43.761	1.00	33.33
ATOM	839	C	ARG	A	120	16.327	21.045	44.699	1.00	33.26
ATOM	840	O	ARG	A	120	16.523	21.138	45.910	1.00	33.07
ATOM	841	CB	ARG	A	120	18.270	22.075	43.592	1.00	33.37
ATOM	842	CG	ARG	A	120	18.759	20.889	42.771	1.00	32.70
ATOM	843	CD	ARG	A	120	20.277	20.732	42.743	1.00	31.64
ATOM	844	NE	ARG	A	120	20.892	21.165	43.992	1.00	31.16
ATOM	845	CZ	ARG	A	120	21.233	20.357	44.993	1.00	31.33
ATOM	846	NH1	ARG	A	120	21.027	19.047	44.920	1.00	30.77
ATOM	847	NH2	ARG	A	120	21.789	20.867	46.077	1.00	31.01
ATOM	848	N	GLU	A	121	15.735	20.001	44.137	1.00	32.98
ATOM	849	CA	GLU	A	121	15.379	18.827	44.904	1.00	32.85
ATOM	850	C	GLU	A	121	15.846	17.630	44.095	1.00	32.17
ATOM	851	O	GLU	A	121	15.744	17.612	42.864	1.00	32.36
ATOM	852	CB	GLU	A	121	13.874	18.767	45.179	1.00	33.21
ATOM	853	CG	GLU	A	121	13.420	17.468	45.826	1.00	35.39
ATOM	854	CD	GLU	A	121	12.189	17.629	46.699	1.00	38.11
ATOM	855	OE1	GLU	A	121	12.326	18.152	47.830	1.00	41.26
ATOM	856	OE2	GLU	A	121	11.091	17.221	46.270	1.00	39.55
ATOM	857	N	GLU	A	122	16.400	16.648	44.786	1.00	31.28
ATOM	858	CA	GLU	A	122	16.886	15.448	44.150	1.00	30.61
ATOM	859	C	GLU	A	122	15.814	14.392	44.298	1.00	30.31
ATOM	860	O	GLU	A	122	15.328	14.165	45.395	1.00	30.07
ATOM	861	CB	GLU	A	122	18.163	14.980	44.833	1.00	30.52
ATOM	862	CG	GLU	A	122	19.293	15.983	44.771	1.00	29.67
ATOM	863	CD	GLU	A	122	19.747	16.253	43.348	1.00	29.79
ATOM	864	OE1	GLU	A	122	20.120	15.280	42.667	1.00	28.28
ATOM	865	OE2	GLU	A	122	19.734	17.439	42.914	1.00	27.93

C
N
C
C
O
C
N
C
C
O
C
C
N
C
C
O
C
C
O
N
C
C
O
C
C
N
N
N
C
C
C
O
C
C
O
O
N
C
C
C
O
O

ATOM	866	N	MET	A	123	15.432	13.751	43.202	1.00	29.74	N
ATOM	867	CA	MET	A	123	14.407	12.728	43.277	1.00	29.77	C
ATOM	868	C	MET	A	123	14.594	11.683	42.202	1.00	29.31	C
ATOM	869	O	MET	A	123	15.408	11.842	41.294	1.00	29.42	O
ATOM	870	CB	MET	A	123	13.016	13.354	43.165	1.00	29.81	C
ATOM	871	CG	MET	A	123	12.749	14.057	41.865	1.00	30.27	C
ATOM	872	SD	MET	A	123	11.181	14.960	41.871	1.00	31.65	S
ATOM	873	CE	MET	A	123	11.582	16.371	42.804	1.00	31.47	C
ATOM	874	N	LYS	A	124	13.835	10.604	42.332	1.00	28.82	N
ATOM	875	CA	LYS	A	124	13.841	9.535	41.352	1.00	28.33	C
ATOM	876	C	LYS	A	124	12.959	9.976	40.191	1.00	27.51	C
ATOM	877	O	LYS	A	124	12.077	10.817	40.361	1.00	26.52	O
ATOM	878	CB	LYS	A	124	13.330	8.242	41.984	1.00	28.45	C
ATOM	879	CG	LYS	A	124	14.153	7.783	43.204	1.00	29.42	C
ATOM	880	CD	LYS	A	124	15.597	7.442	42.804	1.00	30.46	C
ATOM	881	CE	LYS	A	124	16.479	7.067	43.992	1.00	30.52	C
ATOM	882	NZ	LYS	A	124	17.928	7.324	43.699	1.00	30.05	N
ATOM	883	N	PHE	A	125	13.188	9.407	39.014	1.00	26.98	N
ATOM	884	CA	PHE	A	125	12.477	9.871	37.826	1.00	26.83	C
ATOM	885	C	PHE	A	125	10.974	9.762	37.972	1.00	26.77	C
ATOM	886	O	PHE	A	125	10.245	10.673	37.585	1.00	26.38	O
ATOM	887	CB	PHE	A	125	12.927	9.138	36.568	1.00	26.71	C
ATOM	888	CG	PHE	A	125	12.613	9.891	35.316	1.00	26.32	C
ATOM	889	CD1	PHE	A	125	13.476	10.876	34.849	1.00	26.29	C
ATOM	890	CD2	PHE	A	125	11.438	9.657	34.635	1.00	26.06	C
ATOM	891	CE1	PHE	A	125	13.183	11.594	33.715	1.00	25.70	C
ATOM	892	CE2	PHE	A	125	11.140	10.368	33.483	1.00	26.70	C
ATOM	893	CZ	PHE	A	125	12.018	11.341	33.025	1.00	26.87	C
ATOM	894	N	HIS	A	126	10.527	8.649	38.550	1.00	26.77	N
ATOM	895	CA	HIS	A	126	9.113	8.396	38.753	1.00	26.82	C
ATOM	896	C	HIS	A	126	8.517	9.390	39.734	1.00	26.68	C
ATOM	897	O	HIS	A	126	7.326	9.670	39.687	1.00	26.03	O
ATOM	898	CB	HIS	A	126	8.882	6.951	39.238	1.00	27.07	C
ATOM	899	CG	HIS	A	126	9.028	6.767	40.717	1.00	27.39	C
ATOM	900	ND1	HIS	A	126	7.959	6.836	41.582	1.00	28.48	N
ATOM	901	CD2	HIS	A	126	10.115	6.505	41.484	1.00	28.33	C
ATOM	902	CE1	HIS	A	126	8.382	6.636	42.819	1.00	28.68	C
ATOM	903	NE2	HIS	A	126	9.687	6.439	42.788	1.00	27.97	N
ATOM	904	N	GLU	A	127	9.342	9.918	40.630	1.00	27.00	N
ATOM	905	CA	GLU	A	127	8.876	10.930	41.572	1.00	27.34	C
ATOM	906	C	GLU	A	127	8.683	12.259	40.838	1.00	27.36	C
ATOM	907	O	GLU	A	127	7.761	13.026	41.126	1.00	26.86	O
ATOM	908	CB	GLU	A	127	9.862	11.084	42.728	1.00	27.31	C
ATOM	909	CG	GLU	A	127	10.018	9.821	43.552	1.00	28.69	C
ATOM	910	CD	GLU	A	127	10.959	9.992	44.722	1.00	29.65	C
ATOM	911	OE1	GLU	A	127	12.139	10.318	44.485	1.00	29.56	O
ATOM	912	OE2	GLU	A	127	10.509	9.787	45.876	1.00	30.92	O
ATOM	913	N	PHE	A	128	9.554	12.519	39.876	1.00	27.91	N
ATOM	914	CA	PHE	A	128	9.453	13.736	39.068	1.00	28.21	C
ATOM	915	C	PHE	A	128	8.149	13.658	38.282	1.00	29.04	C
ATOM	916	O	PHE	A	128	7.354	14.600	38.255	1.00	29.03	O
ATOM	917	CB	PHE	A	128	10.643	13.838	38.115	1.00	27.74	C
ATOM	918	CG	PHE	A	128	10.427	14.792	36.959	1.00	27.06	C
ATOM	919	CD1	PHE	A	128	10.189	16.138	37.185	1.00	25.41	C
ATOM	920	CD2	PHE	A	128	10.480	14.343	35.657	1.00	25.60	C
ATOM	921	CE1	PHE	A	128	9.985	17.000	36.144	1.00	25.41	C
ATOM	922	CE2	PHE	A	128	10.281	15.215	34.604	1.00	26.59	C
ATOM	923	CZ	PHE	A	128	10.032	16.540	34.846	1.00	25.61	C
ATOM	924	N	VAL	A	129	7.925	12.500	37.677	1.00	29.90	N
ATOM	925	CA	VAL	A	129	6.756	12.285	36.847	1.00	30.82	C
ATOM	926	C	VAL	A	129	5.476	12.447	37.669	1.00	31.61	C

ATOM	927	O	VAL A 129	4.515	13.091	37.234	1.00	31.39	O
ATOM	928	CB	VAL A 129	6.793	10.883	36.213	1.00	30.92	C
ATOM	929	CG1	VAL A 129	5.479	10.582	35.503	1.00	31.15	C
ATOM	930	CG2	VAL A 129	7.975	10.751	35.253	1.00	30.61	C
ATOM	931	N	GLU A 130	5.475	11.858	38.861	1.00	32.42	N
ATOM	932	CA	GLU A 130	4.332	11.946	39.761	1.00	33.13	C
ATOM	933	C	GLU A 130	4.070	13.403	40.126	1.00	33.42	C
ATOM	934	O	GLU A 130	2.930	13.867	40.081	1.00	33.01	O
ATOM	935	CB	GLU A 130	4.587	11.108	41.017	1.00	33.20	C
ATOM	936	CG	GLU A 130	4.537	9.609	40.755	1.00	34.22	C
ATOM	937	CD	GLU A 130	5.294	8.792	41.789	1.00	35.55	C
ATOM	938	OE1	GLU A 130	5.630	9.339	42.861	1.00	37.02	C
ATOM	939	OE2	GLU A 130	5.558	7.598	41.525	1.00	35.99	O
ATOM	940	N	LYS A 131	5.128	14.120	40.486	1.00	34.00	N
ATOM	941	CA	LYS A 131	4.994	15.538	40.800	1.00	34.79	C
ATOM	942	C	LYS A 131	4.354	16.286	39.637	1.00	35.25	C
ATOM	943	O	LYS A 131	3.449	17.089	39.835	1.00	34.81	O
ATOM	944	CB	LYS A 131	6.345	16.171	41.100	1.00	34.89	C
ATOM	945	CG	LYS A 131	6.597	16.477	42.554	1.00	35.57	C
ATOM	946	CD	LYS A 131	7.191	17.868	42.694	1.00	36.25	C
ATOM	947	CE	LYS A 131	7.862	18.072	44.034	1.00	37.30	C
ATOM	948	NZ	LYS A 131	8.108	19.526	44.332	1.00	37.49	N
ATOM	949	N	LEU A 132	4.829	16.038	38.422	1.00	36.00	N
ATOM	950	CA	LEU A 132	4.243	16.701	37.268	1.00	37.05	C
ATOM	951	C	LEU A 132	2.755	16.406	37.170	1.00	37.67	C
ATOM	952	O	LEU A 132	1.963	17.300	36.870	1.00	37.77	O
ATOM	953	CB	LEU A 132	4.919	16.260	35.979	1.00	37.39	C
ATOM	954	CG	LEU A 132	6.310	16.803	35.710	1.00	38.29	C
ATOM	955	CD1	LEU A 132	6.783	16.289	34.363	1.00	39.35	C
ATOM	956	CD2	LEU A 132	6.306	18.314	35.721	1.00	39.70	C
ATOM	957	N	GLN A 133	2.386	15.151	37.417	1.00	38.34	N
ATOM	958	CA	GLN A 133	0.989	14.730	37.373	1.00	39.11	C
ATOM	959	C	GLN A 133	0.132	15.434	38.427	1.00	39.73	C
ATOM	960	O	GLN A 133	-0.966	15.897	38.126	1.00	39.61	O
ATOM	961	CB	GLN A 133	0.886	13.213	37.538	1.00	39.05	C
ATOM	962	N	ASP A 134	0.629	15.507	39.658	1.00	40.61	N
ATOM	963	CA	ASP A 134	-0.108	16.149	40.747	1.00	41.70	C
ATOM	964	C	ASP A 134	-0.398	17.626	40.451	1.00	41.68	C
ATOM	965	O	ASP A 134	-1.470	18.137	40.770	1.00	41.28	O
ATOM	966	CB	ASP A 134	0.678	16.030	42.056	1.00	42.32	C
ATOM	967	CG	ASP A 134	-0.165	16.351	43.286	1.00	44.92	C
ATOM	968	OD1	ASP A 134	-1.235	16.991	43.152	1.00	48.33	O
ATOM	969	OD2	ASP A 134	0.164	15.991	44.442	1.00	48.64	O
ATOM	970	N	ILE A 135	0.563	18.312	39.842	1.00	41.92	N
ATOM	971	CA	ILE A 135	0.383	19.717	39.512	1.00	42.18	C
ATOM	972	C	ILE A 135	-0.715	19.862	38.471	1.00	42.41	C
ATOM	973	O	ILE A 135	-1.635	20.664	38.634	1.00	42.49	O
ATOM	974	CB	ILE A 135	1.696	20.324	38.997	1.00	42.17	C
ATOM	975	CG1	ILE A 135	2.720	20.399	40.131	1.00	42.25	C
ATOM	976	CG2	ILE A 135	1.451	21.708	38.437	1.00	42.11	C
ATOM	977	CD1	ILE A 135	4.151	20.438	39.653	1.00	42.91	C
ATOM	978	N	GLN A 136	-0.614	19.075	37.407	1.00	42.71	N
ATOM	979	CA	GLN A 136	-1.593	19.114	36.333	1.00	43.09	C
ATOM	980	C	GLN A 136	-2.991	18.935	36.900	1.00	43.56	C
ATOM	981	O	GLN A 136	-3.863	19.785	36.707	1.00	43.93	O
ATOM	982	CB	GLN A 136	-1.298	18.031	35.317	1.00	43.10	C
ATOM	983	N	GLN A 137	-3.181	17.844	37.635	1.00	43.82	N
ATOM	984	CA	GLN A 137	-4.486	17.493	38.182	1.00	44.02	C
ATOM	985	C	GLN A 137	-5.023	18.540	39.144	1.00	44.08	C
ATOM	986	O	GLN A 137	-6.202	18.891	39.080	1.00	44.46	O
ATOM	987	CB	GLN A 137	-4.425	16.124	38.868	1.00	44.00	C

ATOM	988	N	ARG A 138	-4.169	19.043	40.031	1.00	43.93	N
ATOM	989	CA	ARG A 138	-4.606	20.022	41.024	1.00	43.76	C
ATOM	990	C	ARG A 138	-4.578	21.445	40.463	1.00	43.34	C
ATOM	991	O	ARG A 138	-4.681	22.415	41.214	1.00	43.54	O
ATOM	992	CB	ARG A 138	-3.751	19.919	42.299	1.00	43.84	C
ATOM	993	CG	ARG A 138	-2.420	20.679	42.279	1.00	44.85	C
ATOM	994	CD	ARG A 138	-1.493	20.279	43.420	1.00	45.82	C
ATOM	995	NE	ARG A 138	-0.372	21.197	43.620	1.00	46.44	N
ATOM	996	CZ	ARG A 138	0.914	20.863	43.499	1.00	47.95	C
ATOM	997	NH1	ARG A 138	1.262	19.627	43.162	1.00	49.02	N
ATOM	998	NH2	ARG A 138	1.864	21.767	43.709	1.00	47.70	N
ATOM	999	N	GLY A 139	-4.450	21.569	39.143	1.00	42.62	N
ATOM	1000	CA	GLY A 139	-4.371	22.871	38.504	1.00	41.93	C
ATOM	1001	C	GLY A 139	-3.389	23.820	39.174	1.00	41.28	C
ATOM	1002	O	GLY A 139	-3.607	25.030	39.196	1.00	41.58	O
ATOM	1003	N	GLY A 140	-2.291	23.292	39.703	1.00	40.31	N
ATOM	1004	CA	GLY A 140	-1.329	24.119	40.410	1.00	39.43	C
ATOM	1005	C	GLY A 140	-0.563	25.070	39.512	1.00	38.71	C
ATOM	1006	O	GLY A 140	-0.495	24.871	38.294	1.00	38.40	O
ATOM	1007	N	GLU A 141	0.003	26.117	40.110	1.00	37.82	N
ATOM	1008	CA	GLU A 141	0.829	27.066	39.363	1.00	37.20	C
ATOM	1009	C	GLU A 141	2.320	26.726	39.491	1.00	35.94	C
ATOM	1010	O	GLU A 141	3.160	27.356	38.845	1.00	35.76	O
ATOM	1011	CB	GLU A 141	0.590	28.510	39.828	1.00	37.63	C
ATOM	1012	CG	GLU A 141	-0.729	29.146	39.379	1.00	39.51	C
ATOM	1013	CD	GLU A 141	-0.936	29.164	37.866	1.00	41.63	C
ATOM	1014	OE1	GLU A 141	0.056	29.147	37.102	1.00	42.42	O
ATOM	1015	OE2	GLU A 141	-2.115	29.202	37.434	1.00	43.77	O
ATOM	1016	N	GLU A 142	2.652	25.742	40.326	1.00	34.29	N
ATOM	1017	CA	GLU A 142	4.045	25.351	40.510	1.00	33.08	C
ATOM	1018	C	GLU A 142	4.655	24.938	39.170	1.00	31.87	C
ATOM	1019	O	GLU A 142	3.958	24.443	38.296	1.00	31.87	O
ATOM	1020	CB	GLU A 142	4.170	24.201	41.519	1.00	32.96	C
ATOM	1021	CG	GLU A 142	5.610	23.914	41.942	1.00	32.71	C
ATOM	1022	CD	GLU A 142	5.761	22.730	42.896	1.00	33.55	C
ATOM	1023	OE1	GLU A 142	4.778	22.001	43.146	1.00	32.41	O
ATOM	1024	OE2	GLU A 142	6.888	22.520	43.398	1.00	34.13	O
ATOM	1025	N	ARG A 143	5.954	25.156	39.017	1.00	30.56	N
ATOM	1026	CA	ARG A 143	6.662	24.766	37.803	1.00	29.79	C
ATOM	1027	C	ARG A 143	7.878	23.939	38.142	1.00	28.70	C
ATOM	1028	O	ARG A 143	8.565	24.210	39.127	1.00	28.88	O
ATOM	1029	CB	ARG A 143	7.142	25.991	37.040	1.00	29.90	C
ATOM	1030	CG	ARG A 143	6.043	26.832	36.441	1.00	30.80	C
ATOM	1031	CD	ARG A 143	6.591	27.992	35.636	1.00	30.70	C
ATOM	1032	NE	ARG A 143	5.538	28.715	34.935	1.00	30.86	N
ATOM	1033	CZ	ARG A 143	5.168	28.487	33.682	1.00	29.53	C
ATOM	1034	NH1	ARG A 143	5.763	27.547	32.941	1.00	27.51	N
ATOM	1035	NH2	ARG A 143	4.196	29.216	33.168	1.00	30.04	N
ATOM	1036	N	LEU A 144	8.156	22.943	37.315	1.00	27.27	N
ATOM	1037	CA	LEU A 144	9.319	22.102	37.518	1.00	26.47	C
ATOM	1038	C	LEU A 144	10.220	22.148	36.297	1.00	25.40	C
ATOM	1039	O	LEU A 144	9.755	22.330	35.172	1.00	25.39	O
ATOM	1040	CB	LEU A 144	8.893	20.652	37.753	1.00	26.70	C
ATOM	1041	CG	LEU A 144	7.922	20.416	38.915	1.00	27.15	C
ATOM	1042	CD1	LEU A 144	7.575	18.921	39.059	1.00	27.40	C
ATOM	1043	CD2	LEU A 144	8.488	20.956	40.221	1.00	27.42	C
ATOM	1044	N	TYR A 145	11.511	21.974	36.527	1.00	23.90	N
ATOM	1045	CA	TYR A 145	12.458	21.871	35.441	1.00	23.15	C
ATOM	1046	C	TYR A 145	13.516	20.857	35.868	1.00	23.04	C
ATOM	1047	O	TYR A 145	14.328	21.130	36.740	1.00	22.58	O
ATOM	1048	CB	TYR A 145	13.080	23.235	35.107	1.00	23.01	C

ATOM	1049	CG	TYR	A	145	13.522	23.419	33.666	1.00	21.34	C
ATOM	1050	CD1	TYR	A	145	13.824	22.333	32.863	1.00	20.11	C
ATOM	1051	CD2	TYR	A	145	13.662	24.691	33.119	1.00	21.14	C
ATOM	1052	CE1	TYR	A	145	14.219	22.494	31.563	1.00	20.22	C
ATOM	1053	CE2	TYR	A	145	14.075	24.866	31.807	1.00	19.81	C
ATOM	1054	CZ	TYR	A	145	14.349	23.764	31.027	1.00	19.19	C
ATOM	1055	OH	TYR	A	145	14.736	23.903	29.699	1.00	18.36	O
ATOM	1056	N	LEU	A	146	13.461	19.666	35.287	1.00	22.75	N
ATOM	1057	CA	LEU	A	146	14.490	18.661	35.516	1.00	23.17	C
ATOM	1058	C	LEU	A	146	15.720	19.005	34.679	1.00	22.99	C
ATOM	1059	O	LEU	A	146	15.604	19.264	33.490	1.00	22.81	O
ATOM	1060	CB	LEU	A	146	13.980	17.282	35.115	1.00	23.33	C
ATOM	1061	CG	LEU	A	146	14.992	16.141	35.274	1.00	24.98	C
ATOM	1062	CD1	LEU	A	146	14.276	14.846	35.622	1.00	24.56	C
ATOM	1063	CD2	LEU	A	146	15.849	15.946	34.016	1.00	26.13	C
ATOM	1064	N	GLN	A	147	16.891	18.983	35.299	1.00	23.05	N
ATOM	1065	CA	GLN	A	147	18.135	19.314	34.619	1.00	23.65	C
ATOM	1066	C	GLN	A	147	19.158	18.394	35.231	1.00	23.86	C
ATOM	1067	O	GLN	A	147	19.573	18.597	36.364	1.00	24.61	O
ATOM	1068	CB	GLN	A	147	18.516	20.793	34.821	1.00	23.53	C
ATOM	1069	CG	GLN	A	147	17.386	21.770	34.461	1.00	24.34	C
ATOM	1070	CD	GLN	A	147	17.800	23.238	34.482	1.00	26.20	C
ATOM	1071	OE1	GLN	A	147	17.034	24.114	34.035	1.00	29.67	O
ATOM	1072	NE2	GLN	A	147	18.979	23.514	34.988	1.00	21.33	N
ATOM	1073	N	GLN	A	148	19.542	17.369	34.485	1.00	24.35	N
ATOM	1074	CA	GLN	A	148	20.393	16.314	35.006	1.00	24.70	C
ATOM	1075	C	GLN	A	148	21.319	15.753	33.964	1.00	25.16	C
ATOM	1076	O	GLN	A	148	20.898	15.378	32.866	1.00	24.21	O
ATOM	1077	CB	GLN	A	148	19.525	15.172	35.526	1.00	24.97	C
ATOM	1078	CG	GLN	A	148	20.317	13.940	35.953	1.00	25.54	C
ATOM	1079	CD	GLN	A	148	21.275	14.256	37.085	1.00	26.98	C
ATOM	1080	OE1	GLN	A	148	20.892	14.941	38.042	1.00	26.73	O
ATOM	1081	NE2	GLN	A	148	22.522	13.786	36.976	1.00	26.37	N
ATOM	1082	N	THR	A	149	22.592	15.704	34.321	1.00	25.94	N
ATOM	1083	CA	THR	A	149	23.603	15.134	33.466	1.00	27.43	C
ATOM	1084	C	THR	A	149	23.369	13.632	33.324	1.00	27.29	C
ATOM	1085	O	THR	A	149	23.081	12.965	34.303	1.00	27.40	O
ATOM	1086	CB	THR	A	149	24.990	15.430	34.075	1.00	27.91	C
ATOM	1087	OG1	THR	A	149	25.282	16.829	33.901	1.00	30.41	O
ATOM	1088	CG2	THR	A	149	26.078	14.776	33.276	1.00	30.12	C
ATOM	1089	N	LEU	A	150	23.461	13.117	32.100	1.00	27.79	N
ATOM	1090	CA	LEU	A	150	23.321	11.690	31.832	1.00	28.02	C
ATOM	1091	C	LEU	A	150	24.549	10.935	32.364	1.00	28.42	C
ATOM	1092	O	LEU	A	150	25.682	11.261	32.002	1.00	28.86	O
ATOM	1093	CB	LEU	A	150	23.194	11.435	30.326	1.00	28.08	C
ATOM	1094	CG	LEU	A	150	21.929	11.937	29.622	1.00	28.70	C
ATOM	1095	CD1	LEU	A	150	22.016	11.725	28.117	1.00	28.62	C
ATOM	1096	CD2	LEU	A	150	20.697	11.260	30.175	1.00	29.47	C
ATOM	1097	N	ASN	A	151	24.332	9.928	33.203	1.00	28.58	N
ATOM	1098	CA	ASN	A	151	25.434	9.156	33.781	1.00	28.79	C
ATOM	1099	C	ASN	A	151	25.194	7.636	33.782	1.00	29.24	C
ATOM	1100	O	ASN	A	151	24.197	7.163	33.238	1.00	29.60	O
ATOM	1101	CB	ASN	A	151	25.657	9.631	35.208	1.00	28.77	C
ATOM	1102	CG	ASN	A	151	24.459	9.367	36.084	1.00	28.05	C
ATOM	1103	OD1	ASN	A	151	23.936	8.246	36.126	1.00	29.02	O
ATOM	1104	ND2	ASN	A	151	24.000	10.396	36.772	1.00	26.99	N
ATOM	1105	N	ASP	A	152	26.083	6.885	34.437	1.00	29.71	N
ATOM	1106	CA	ASP	A	152	26.039	5.410	34.448	1.00	30.06	C
ATOM	1107	C	ASP	A	152	24.850	4.733	35.075	1.00	29.51	C
ATOM	1108	O	ASP	A	152	24.771	3.503	35.026	1.00	28.98	O
ATOM	1109	CB	ASP	A	152	27.199	4.829	35.250	1.00	30.99	C

ATOM	1110	CG	ASP	A	152	28.447	5.561	35.048	1.00	33.80
ATOM	1111	OD1	ASP	A	152	28.636	6.071	33.918	1.00	40.13
ATOM	1112	OD2	ASP	A	152	29.274	5.719	35.960	1.00	35.65
ATOM	1113	N	THR	A	153	23.959	5.468	35.722	1.00	28.96
ATOM	1114	CA	THR	A	153	22.831	4.792	36.359	1.00	28.42
ATOM	1115	C	THR	A	153	21.685	4.594	35.387	1.00	27.75
ATOM	1116	O	THR	A	153	20.730	3.909	35.712	1.00	27.68
ATOM	1117	CB	THR	A	153	22.330	5.552	37.584	1.00	28.53
ATOM	1118	OG1	THR	A	153	21.833	6.836	37.193	1.00	28.91
ATOM	1119	CG2	THR	A	153	23.473	5.855	38.540	1.00	29.18
ATOM	1120	N	VAL	A	154	21.766	5.171	34.194	1.00	26.76
ATOM	1121	CA	VAL	A	154	20.671	4.996	33.246	1.00	26.61
ATOM	1122	C	VAL	A	154	20.583	3.531	32.895	1.00	26.77
ATOM	1123	O	VAL	A	154	21.592	2.832	32.923	1.00	27.36
ATOM	1124	CB	VAL	A	154	20.838	5.826	31.964	1.00	26.05
ATOM	1125	CG1	VAL	A	154	20.914	7.279	32.312	1.00	26.14
ATOM	1126	CG2	VAL	A	154	22.071	5.383	31.182	1.00	25.62
ATOM	1127	N	GLY	A	155	19.379	3.072	32.568	1.00	26.71
ATOM	1128	CA	GLY	A	155	19.147	1.674	32.252	1.00	26.42
ATOM	1129	C	GLY	A	155	19.531	1.266	30.840	1.00	26.67
ATOM	1130	O	GLY	A	155	19.894	2.093	29.983	1.00	26.52
ATOM	1131	N	ARG	A	156	19.390	-0.028	30.599	1.00	26.49
ATOM	1132	CA	ARG	A	156	19.811	-0.684	29.364	1.00	26.54
ATOM	1133	C	ARG	A	156	19.305	-0.053	28.068	1.00	25.95
ATOM	1134	O	ARG	A	156	20.089	0.194	27.160	1.00	26.26
ATOM	1135	CB	ARG	A	156	19.429	-2.165	29.418	1.00	26.28
ATOM	1136	N	LYS	A	157	18.004	0.175	27.964	1.00	25.43
ATOM	1137	CA	LYS	A	157	17.460	0.777	26.756	1.00	25.11
ATOM	1138	C	LYS	A	157	18.026	2.198	26.517	1.00	25.50
ATOM	1139	O	LYS	A	157	18.249	2.595	25.372	1.00	25.14
ATOM	1140	CB	LYS	A	157	15.927	0.796	26.809	1.00	24.88
ATOM	1141	CG	LYS	A	157	15.255	-0.572	26.516	1.00	24.25
ATOM	1142	N	ILE	A	158	18.257	2.956	27.585	1.00	25.35
ATOM	1143	CA	ILE	A	158	18.766	4.326	27.432	1.00	25.61
ATOM	1144	C	ILE	A	158	20.205	4.250	26.971	1.00	25.76
ATOM	1145	O	ILE	A	158	20.661	5.058	26.166	1.00	25.39
ATOM	1146	CB	ILE	A	158	18.662	5.107	28.740	1.00	25.41
ATOM	1147	CG1	ILE	A	158	17.202	5.245	29.162	1.00	25.92
ATOM	1148	CG2	ILE	A	158	19.297	6.475	28.601	1.00	25.61
ATOM	1149	CD1	ILE	A	158	16.331	5.924	28.174	1.00	27.88
ATOM	1150	N	VAL	A	159	20.909	3.251	27.481	1.00	26.10
ATOM	1151	CA	VAL	A	159	22.276	3.010	27.079	1.00	26.57
ATOM	1152	C	VAL	A	159	22.279	2.703	25.585	1.00	26.40
ATOM	1153	O	VAL	A	159	23.074	3.248	24.840	1.00	26.32
ATOM	1154	CB	VAL	A	159	22.895	1.850	27.883	1.00	26.94
ATOM	1155	CG1	VAL	A	159	24.136	1.310	27.193	1.00	27.73
ATOM	1156	CG2	VAL	A	159	23.223	2.308	29.298	1.00	26.73
ATOM	1157	N	MET	A	160	21.363	1.852	25.145	1.00	26.48
ATOM	1158	CA	MET	A	160	21.270	1.518	23.721	1.00	26.52
ATOM	1159	C	MET	A	160	20.985	2.771	22.880	1.00	25.02
ATOM	1160	O	MET	A	160	21.600	2.986	21.845	1.00	24.31
ATOM	1161	CB	MET	A	160	20.183	0.476	23.485	1.00	27.05
ATOM	1162	CG	MET	A	160	20.540	-0.900	24.001	1.00	30.45
ATOM	1163	SD	MET	A	160	21.843	-1.730	23.058	1.00	34.87
ATOM	1164	CE	MET	A	160	20.957	-2.028	21.496	1.00	36.94
ATOM	1165	N	ASP	A	161	20.047	3.589	23.342	1.00	23.96
ATOM	1166	CA	ASP	A	161	19.699	4.831	22.665	1.00	23.14
ATOM	1167	C	ASP	A	161	20.890	5.781	22.550	1.00	21.93
ATOM	1168	O	ASP	A	161	21.167	6.323	21.480	1.00	21.41
ATOM	1169	CB	ASP	A	161	18.549	5.528	23.402	1.00	23.13
ATOM	1170	CG	ASP	A	161	17.250	4.762	23.293	1.00	24.08

C
O
O
N
C
C
O
C
O
C
N
C
C
O
C
C
N
C
C
O
N
C
C
O
C
C
N
C
C
C
C
C
N
C
C
C
C
S
C
N
C
C
O
C
C

ATOM	1171	OD1	ASP	A	161	17.182	3.812	22.478	1.00	21.27	O
ATOM	1172	OD2	ASP	A	161	16.248	5.035	23.980	1.00	25.87	O
ATOM	1173	N	PHE	A	162	21.575	5.973	23.671	1.00	20.59	N
ATOM	1174	CA	PHE	A	162	22.712	6.859	23.765	1.00	19.97	C
ATOM	1175	C	PHE	A	162	23.802	6.409	22.809	1.00	19.19	C
ATOM	1176	O	PHE	A	162	24.410	7.213	22.160	1.00	18.98	O
ATOM	1177	CB	PHE	A	162	23.223	6.859	25.220	1.00	20.44	C
ATOM	1178	CG	PHE	A	162	24.386	7.755	25.470	1.00	20.78	C
ATOM	1179	CD1	PHE	A	162	24.206	9.107	25.673	1.00	25.50	C
ATOM	1180	CD2	PHE	A	162	25.662	7.245	25.542	1.00	23.71	C
ATOM	1181	CE1	PHE	A	162	25.290	9.930	25.935	1.00	25.85	C
ATOM	1182	CE2	PHE	A	162	26.755	8.072	25.795	1.00	24.89	C
ATOM	1183	CZ	PHE	A	162	26.572	9.394	26.001	1.00	23.50	C
ATOM	1184	N	LEU	A	163	24.062	5.115	22.744	1.00	19.25	N
ATOM	1185	CA	LEU	A	163	25.084	4.597	21.838	1.00	19.42	C
ATOM	1186	C	LEU	A	163	24.715	4.804	20.361	1.00	18.63	C
ATOM	1187	O	LEU	A	163	25.585	4.831	19.493	1.00	18.21	O
ATOM	1188	CB	LEU	A	163	25.297	3.115	22.104	1.00	19.49	C
ATOM	1189	CG	LEU	A	163	25.988	2.812	23.422	1.00	20.88	C
ATOM	1190	CD1	LEU	A	163	25.980	1.319	23.651	1.00	22.44	C
ATOM	1191	CD2	LEU	A	163	27.407	3.368	23.396	1.00	21.73	C
ATOM	1192	N	GLY	A	164	23.419	4.917	20.104	1.00	18.01	N
ATOM	1193	CA	GLY	A	164	22.889	5.158	18.779	1.00	18.55	C
ATOM	1194	C	GLY	A	164	22.873	6.622	18.355	1.00	18.40	C
ATOM	1195	O	GLY	A	164	22.406	6.921	17.256	1.00	19.24	O
ATOM	1196	N	PHE	A	165	23.365	7.521	19.209	1.00	17.50	N
ATOM	1197	CA	PHE	A	165	23.493	8.913	18.831	1.00	17.67	C
ATOM	1198	C	PHE	A	165	24.497	8.955	17.663	1.00	17.49	C
ATOM	1199	O	PHE	A	165	25.293	8.028	17.497	1.00	16.10	O
ATOM	1200	CB	PHE	A	165	23.984	9.753	20.018	1.00	17.47	C
ATOM	1201	CG	PHE	A	165	22.932	10.004	21.098	1.00	18.51	C
ATOM	1202	CD1	PHE	A	165	21.645	9.479	21.009	1.00	19.26	C
ATOM	1203	CD2	PHE	A	165	23.242	10.784	22.203	1.00	18.52	C
ATOM	1204	CE1	PHE	A	165	20.697	9.730	21.996	1.00	18.44	C
ATOM	1205	CE2	PHE	A	165	22.303	11.030	23.199	1.00	18.52	C
ATOM	1206	CZ	PHE	A	165	21.032	10.500	23.100	1.00	18.90	C
ATOM	1207	N	ASN	A	166	24.466	10.009	16.854	1.00	17.77	N
ATOM	1208	CA	ASN	A	166	25.393	10.110	15.712	1.00	18.29	C
ATOM	1209	C	ASN	A	166	26.787	10.622	16.129	1.00	18.73	C
ATOM	1210	O	ASN	A	166	27.156	11.795	15.897	1.00	19.54	O
ATOM	1211	CB	ASN	A	166	24.793	10.972	14.598	1.00	17.92	C
ATOM	1212	CG	ASN	A	166	25.571	10.861	13.293	1.00	17.28	C
ATOM	1213	OD1	ASN	A	166	26.679	10.289	13.262	1.00	16.53	O
ATOM	1214	ND2	ASN	A	166	24.994	11.395	12.204	1.00	12.92	N
ATOM	1215	N	TRP	A	167	27.527	9.739	16.789	1.00	19.26	N
ATOM	1216	CA	TRP	A	167	28.867	10.035	17.264	1.00	19.76	C
ATOM	1217	C	TRP	A	167	29.785	10.266	16.084	1.00	19.88	C
ATOM	1218	O	TRP	A	167	30.731	11.031	16.169	1.00	19.35	O
ATOM	1219	CB	TRP	A	167	29.384	8.864	18.130	1.00	19.92	C
ATOM	1220	CG	TRP	A	167	28.556	8.728	19.351	1.00	20.39	C
ATOM	1221	CD1	TRP	A	167	27.686	7.727	19.656	1.00	20.85	C
ATOM	1222	CD2	TRP	A	167	28.445	9.686	20.400	1.00	20.99	C
ATOM	1223	NE1	TRP	A	167	27.059	7.995	20.851	1.00	21.18	N
ATOM	1224	CE2	TRP	A	167	27.509	9.194	21.325	1.00	20.70	C
ATOM	1225	CE3	TRP	A	167	29.065	10.916	20.664	1.00	21.55	C
ATOM	1226	CZ2	TRP	A	167	27.183	9.871	22.488	1.00	22.63	C
ATOM	1227	CZ3	TRP	A	167	28.731	11.589	21.804	1.00	21.78	C
ATOM	1228	CH2	TRP	A	167	27.789	11.071	22.706	1.00	22.79	C
ATOM	1229	N	ASN	A	168	29.529	9.577	14.979	1.00	20.07	N
ATOM	1230	CA	ASN	A	168	30.374	9.759	13.818	1.00	20.50	C
ATOM	1231	C	ASN	A	168	30.396	11.237	13.421	1.00	20.67	C

ATOM	1232	O	ASN A 168	31.465	11.806	13.207	1.00	19.54	O
ATOM	1233	CB	ASN A 168	29.917	8.930	12.628	1.00	20.87	C
ATOM	1234	CG	ASN A 168	30.818	9.129	11.423	1.00	22.96	C
ATOM	1235	OD1	ASN A 168	32.027	8.923	11.522	1.00	25.54	O
ATOM	1236	ND2	ASN A 168	30.247	9.578	10.295	1.00	23.09	N
ATOM	1237	N	TRP A 169	29.211	11.844	13.338	1.00	20.57	N
ATOM	1238	CA	TRP A 169	29.106	13.226	12.917	1.00	20.23	C
ATOM	1239	C	TRP A 169	29.653	14.188	13.972	1.00	20.59	C
ATOM	1240	O	TRP A 169	30.367	15.118	13.634	1.00	20.09	O
ATOM	1241	CB	TRP A 169	27.662	13.618	12.570	1.00	20.35	C
ATOM	1242	CG	TRP A 169	27.542	15.101	12.238	1.00	19.49	C
ATOM	1243	CD1	TRP A 169	27.769	15.693	11.026	1.00	19.03	C
ATOM	1244	CD2	TRP A 169	27.203	16.157	13.137	1.00	20.25	C
ATOM	1245	NE1	TRP A 169	27.578	17.052	11.117	1.00	20.36	N
ATOM	1246	CE2	TRP A 169	27.244	17.366	12.406	1.00	19.45	C
ATOM	1247	CE3	TRP A 169	26.874	16.207	14.492	1.00	20.60	C
ATOM	1248	CZ2	TRP A 169	26.964	18.600	12.975	1.00	21.03	C
ATOM	1249	CZ3	TRP A 169	26.614	17.433	15.064	1.00	23.09	C
ATOM	1250	CH2	TRP A 169	26.649	18.621	14.297	1.00	23.61	C
ATOM	1251	N	ILE A 170	29.326	13.979	15.239	1.00	20.47	N
ATOM	1252	CA	ILE A 170	29.759	14.926	16.241	1.00	20.67	C
ATOM	1253	C	ILE A 170	31.262	14.772	16.567	1.00	21.18	C
ATOM	1254	O	ILE A 170	31.943	15.758	16.836	1.00	21.20	O
ATOM	1255	CB	ILE A 170	28.842	14.892	17.483	1.00	20.61	C
ATOM	1256	CG1	ILE A 170	28.900	16.231	18.221	1.00	20.05	C
ATOM	1257	CG2	ILE A 170	29.191	13.755	18.402	1.00	19.89	C
ATOM	1258	CD1	ILE A 170	27.865	16.353	19.329	1.00	21.00	C
ATOM	1259	N	ASN A 171	31.780	13.556	16.527	1.00	20.74	N
ATOM	1260	CA	ASN A 171	33.214	13.355	16.715	1.00	21.80	C
ATOM	1261	C	ASN A 171	34.024	14.093	15.634	1.00	22.34	C
ATOM	1262	O	ASN A 171	35.093	14.652	15.916	1.00	21.98	O
ATOM	1263	CB	ASN A 171	33.581	11.857	16.718	1.00	21.55	C
ATOM	1264	CG	ASN A 171	33.111	11.124	17.981	1.00	21.20	C
ATOM	1265	OD1	ASN A 171	32.637	11.724	18.963	1.00	21.35	O
ATOM	1266	ND2	ASN A 171	33.263	9.830	17.962	1.00	19.36	N
ATOM	1267	N	LYS A 172	33.529	14.097	14.400	1.00	22.97	N
ATOM	1268	CA	LYS A 172	34.218	14.847	13.353	1.00	24.09	C
ATOM	1269	C	LYS A 172	34.167	16.342	13.656	1.00	23.61	C
ATOM	1270	O	LYS A 172	35.166	17.041	13.511	1.00	23.54	O
ATOM	1271	CB	LYS A 172	33.669	14.529	11.961	1.00	24.58	C
ATOM	1272	CG	LYS A 172	34.087	13.163	11.491	1.00	27.56	C
ATOM	1273	CD	LYS A 172	33.653	12.852	10.034	1.00	31.03	C
ATOM	1274	CE	LYS A 172	34.323	11.576	9.565	1.00	33.01	C
ATOM	1275	NZ	LYS A 172	34.357	11.406	8.070	1.00	36.89	N
ATOM	1276	N	GLN A 173	33.019	16.834	14.109	1.00	23.74	N
ATOM	1277	CA	GLN A 173	32.914	18.245	14.482	1.00	23.87	C
ATOM	1278	C	GLN A 173	33.960	18.578	15.547	1.00	23.86	C
ATOM	1279	O	GLN A 173	34.739	19.500	15.374	1.00	24.25	O
ATOM	1280	CB	GLN A 173	31.517	18.590	14.993	1.00	23.63	C
ATOM	1281	CG	GLN A 173	30.451	18.619	13.916	1.00	24.53	C
ATOM	1282	CD	GLN A 173	30.753	19.634	12.851	1.00	25.08	C
ATOM	1283	OE1	GLN A 173	31.194	20.740	13.157	1.00	26.69	O
ATOM	1284	NE2	GLN A 173	30.512	19.273	11.601	1.00	23.36	N
ATOM	1285	N	GLN A 174	33.963	17.822	16.635	1.00	23.63	N
ATOM	1286	CA	GLN A 174	34.925	17.995	17.717	1.00	23.91	C
ATOM	1287	C	GLN A 174	36.365	18.053	17.161	1.00	24.01	C
ATOM	1288	O	GLN A 174	37.133	18.962	17.480	1.00	23.39	O
ATOM	1289	CB	GLN A 174	34.783	16.840	18.717	1.00	23.67	C
ATOM	1290	CG	GLN A 174	35.688	16.913	19.934	1.00	24.07	C
ATOM	1291	CD	GLN A 174	35.595	15.668	20.799	1.00	25.12	C
ATOM	1292	OE1	GLN A 174	35.229	14.602	20.312	1.00	25.70	O

ATOM	1293	NE2	GLN	A	174	35.901	15.803	22.084	1.00	23.63	N
ATOM	1294	N	GLY	A	175	36.706	17.080	16.334	1.00	23.75	N
ATOM	1295	CA	GLY	A	175	38.005	17.028	15.696	1.00	24.73	C
ATOM	1296	C	GLY	A	175	38.268	18.209	14.768	1.00	25.21	C
ATOM	1297	O	GLY	A	175	39.310	18.842	14.854	1.00	25.60	O
ATOM	1298	N	LYS	A	176	37.327	18.518	13.888	1.00	25.81	N
ATOM	1299	CA	LYS	A	176	37.491	19.634	12.961	1.00	26.88	C
ATOM	1300	C	LYS	A	176	37.719	20.997	13.638	1.00	26.95	C
ATOM	1301	O	LYS	A	176	38.490	21.803	13.136	1.00	26.74	O
ATOM	1302	CB	LYS	A	176	36.256	19.778	12.077	1.00	27.35	C
ATOM	1303	CG	LYS	A	176	36.093	18.716	11.009	1.00	29.87	C
ATOM	1304	CD	LYS	A	176	34.894	19.085	10.137	1.00	33.21	C
ATOM	1305	CE	LYS	A	176	33.948	17.932	9.961	1.00	35.30	C
ATOM	1306	NZ	LYS	A	176	32.558	18.388	9.643	1.00	37.22	N
ATOM	1307	N	ARG	A	177	37.031	21.257	14.749	1.00	26.93	N
ATOM	1308	CA	ARG	A	177	37.153	22.539	15.435	1.00	27.36	C
ATOM	1309	C	ARG	A	177	38.241	22.611	16.509	1.00	26.63	C
ATOM	1310	O	ARG	A	177	38.402	23.650	17.135	1.00	25.85	O
ATOM	1311	CB	ARG	A	177	35.832	22.902	16.114	1.00	27.86	C
ATOM	1312	CG	ARG	A	177	34.625	22.712	15.267	1.00	29.88	C
ATOM	1313	CD	ARG	A	177	34.653	23.460	13.973	1.00	32.39	C
ATOM	1314	NE	ARG	A	177	33.683	22.854	13.084	1.00	34.54	N
ATOM	1315	CZ	ARG	A	177	33.841	22.701	11.790	1.00	37.31	C
ATOM	1316	NH1	ARG	A	177	34.952	23.110	11.189	1.00	38.34	N
ATOM	1317	NH2	ARG	A	177	32.877	22.126	11.088	1.00	38.88	N
ATOM	1318	N	GLY	A	178	38.950	21.510	16.743	1.00	26.35	N
ATOM	1319	CA	GLY	A	178	39.998	21.470	17.753	1.00	25.46	C
ATOM	1320	C	GLY	A	178	39.473	21.514	19.176	1.00	25.27	C
ATOM	1321	O	GLY	A	178	40.213	21.818	20.123	1.00	25.46	O
ATOM	1322	N	TRP	A	179	38.199	21.202	19.359	1.00	24.30	N
ATOM	1323	CA	TRP	A	179	37.639	21.260	20.692	1.00	24.02	C
ATOM	1324	C	TRP	A	179	38.290	20.266	21.638	1.00	23.85	C
ATOM	1325	O	TRP	A	179	38.958	19.333	21.226	1.00	22.62	O
ATOM	1326	CB	TRP	A	179	36.136	21.011	20.674	1.00	23.73	C
ATOM	1327	CG	TRP	A	179	35.346	22.061	19.962	1.00	23.92	C
ATOM	1328	CD1	TRP	A	179	35.787	23.291	19.531	1.00	22.69	C
ATOM	1329	CD2	TRP	A	179	33.968	21.981	19.594	1.00	23.74	C
ATOM	1330	NE1	TRP	A	179	34.765	23.968	18.912	1.00	24.45	N
ATOM	1331	CE2	TRP	A	179	33.636	23.183	18.929	1.00	24.47	C
ATOM	1332	CE3	TRP	A	179	32.984	21.002	19.730	1.00	23.10	C
ATOM	1333	CZ2	TRP	A	179	32.379	23.422	18.414	1.00	23.69	C
ATOM	1334	CZ3	TRP	A	179	31.733	21.241	19.211	1.00	21.89	C
ATOM	1335	CH2	TRP	A	179	31.435	22.445	18.573	1.00	23.90	C
ATOM	1336	N	GLY	A	180	38.080	20.499	22.925	1.00	24.08	N
ATOM	1337	CA	GLY	A	180	38.488	19.558	23.941	1.00	24.08	C
ATOM	1338	C	GLY	A	180	37.437	18.476	24.103	1.00	24.64	C
ATOM	1339	O	GLY	A	180	36.618	18.234	23.202	1.00	24.66	O
ATOM	1340	N	GLN	A	181	37.437	17.830	25.261	1.00	24.77	N
ATOM	1341	CA	GLN	A	181	36.575	16.682	25.471	1.00	25.23	C
ATOM	1342	C	GLN	A	181	35.117	17.039	25.725	1.00	24.50	C
ATOM	1343	O	GLN	A	181	34.779	18.151	26.149	1.00	24.42	O
ATOM	1344	CB	GLN	A	181	37.094	15.847	26.646	1.00	25.76	C
ATOM	1345	CG	GLN	A	181	36.720	16.409	28.025	1.00	28.85	C
ATOM	1346	CD	GLN	A	181	37.046	15.434	29.156	1.00	33.38	C
ATOM	1347	OE1	GLN	A	181	38.186	15.003	29.293	1.00	36.10	O
ATOM	1348	NE2	GLN	A	181	36.044	15.085	29.958	1.00	36.04	N
ATOM	1349	N	LEU	A	182	34.262	16.078	25.423	1.00	23.71	N
ATOM	1350	CA	LEU	A	182	32.857	16.110	25.792	1.00	23.70	C
ATOM	1351	C	LEU	A	182	32.876	16.040	27.317	1.00	22.68	C
ATOM	1352	O	LEU	A	182	33.406	15.079	27.849	1.00	21.72	O
ATOM	1353	CB	LEU	A	182	32.179	14.836	25.273	1.00	23.82	C

ATOM	1354	CG	LEU	A	182	30.661	14.693	25.199	1.00	26.38	C
ATOM	1355	CD1	LEU	A	182	30.243	13.231	25.368	1.00	26.04	C
ATOM	1356	CD2	LEU	A	182	29.977	15.501	26.192	1.00	29.73	C
ATOM	1357	N	THR	A	183	32.323	17.031	28.021	1.00	21.93	N
ATOM	1358	CA	THR	A	183	32.300	16.971	29.484	1.00	21.44	C
ATOM	1359	C	THR	A	183	31.004	16.388	29.984	1.00	21.27	C
ATOM	1360	O	THR	A	183	30.972	15.766	31.032	1.00	21.29	O
ATOM	1361	CB	THR	A	183	32.490	18.362	30.150	1.00	21.73	C
ATOM	1362	OG1	THR	A	183	31.463	19.257	29.715	1.00	20.35	O
ATOM	1363	CG2	THR	A	183	33.796	19.015	29.725	1.00	20.91	C
ATOM	1364	N	SER	A	184	29.918	16.590	29.251	1.00	21.42	N
ATOM	1365	CA	SER	A	184	28.649	16.053	29.689	1.00	21.45	C
ATOM	1366	C	SER	A	184	27.537	16.272	28.714	1.00	21.46	C
ATOM	1367	O	SER	A	184	27.672	17.031	27.753	1.00	21.17	O
ATOM	1368	CB	SER	A	184	28.243	16.687	31.006	1.00	21.74	C
ATOM	1369	OG	SER	A	184	27.919	18.049	30.845	1.00	23.24	O
ATOM	1370	N	ASN	A	185	26.445	15.575	28.982	1.00	21.21	N
ATOM	1371	CA	ASN	A	185	25.216	15.712	28.245	1.00	22.29	C
ATOM	1372	C	ASN	A	185	24.154	16.005	29.273	1.00	22.08	C
ATOM	1373	O	ASN	A	185	23.886	15.184	30.135	1.00	22.25	O
ATOM	1374	CB	ASN	A	185	24.856	14.416	27.497	1.00	22.67	C
ATOM	1375	CG	ASN	A	185	25.885	14.037	26.454	1.00	23.57	C
ATOM	1376	OD1	ASN	A	185	26.646	13.097	26.655	1.00	27.51	O
ATOM	1377	ND2	ASN	A	185	25.905	14.754	25.329	1.00	24.27	N
ATOM	1378	N	LEU	A	186	23.574	17.189	29.202	1.00	22.01	N
ATOM	1379	CA	LEU	A	186	22.529	17.563	30.135	1.00	22.03	C
ATOM	1380	C	LEU	A	186	21.170	17.219	29.558	1.00	22.00	C
ATOM	1381	O	LEU	A	186	20.844	17.592	28.435	1.00	22.34	O
ATOM	1382	CB	LEU	A	186	22.578	19.065	30.423	1.00	21.68	C
ATOM	1383	CG	LEU	A	186	21.707	19.538	31.588	1.00	22.25	C
ATOM	1384	CD1	LEU	A	186	22.252	18.985	32.891	1.00	22.43	C
ATOM	1385	CD2	LEU	A	186	21.643	21.098	31.648	1.00	23.35	C
ATOM	1386	N	LEU	A	187	20.377	16.518	30.344	1.00	22.19	N
ATOM	1387	CA	LEU	A	187	19.009	16.198	29.979	1.00	21.95	C
ATOM	1388	C	LEU	A	187	18.149	17.286	30.603	1.00	21.91	C
ATOM	1389	O	LEU	A	187	18.232	17.521	31.823	1.00	21.61	O
ATOM	1390	CB	LEU	A	187	18.616	14.830	30.527	1.00	21.55	C
ATOM	1391	CG	LEU	A	187	17.129	14.488	30.477	1.00	22.78	C
ATOM	1392	CD1	LEU	A	187	16.616	14.429	29.054	1.00	23.66	C
ATOM	1393	CD2	LEU	A	187	16.866	13.143	31.174	1.00	22.84	C
ATOM	1394	N	LEU	A	188	17.348	17.956	29.770	1.00	21.56	N
ATOM	1395	CA	LEU	A	188	16.461	19.000	30.227	1.00	22.26	C
ATOM	1396	C	LEU	A	188	14.993	18.662	29.920	1.00	22.55	C
ATOM	1397	O	LEU	A	188	14.588	18.517	28.760	1.00	23.16	O
ATOM	1398	CB	LEU	A	188	16.827	20.327	29.562	1.00	22.84	C
ATOM	1399	CG	LEU	A	188	18.244	20.840	29.821	1.00	22.60	C
ATOM	1400	CD1	LEU	A	188	18.967	21.111	28.523	1.00	23.92	C
ATOM	1401	CD2	LEU	A	188	18.177	22.088	30.655	1.00	24.56	C
ATOM	1402	N	ILE	A	189	14.181	18.577	30.958	1.00	22.15	N
ATOM	1403	CA	ILE	A	189	12.769	18.315	30.756	1.00	22.34	C
ATOM	1404	C	ILE	A	189	11.996	19.395	31.460	1.00	22.37	C
ATOM	1405	O	ILE	A	189	12.072	19.528	32.692	1.00	22.77	O
ATOM	1406	CB	ILE	A	189	12.377	16.953	31.281	1.00	21.60	C
ATOM	1407	CG1	ILE	A	189	13.254	15.888	30.638	1.00	21.85	C
ATOM	1408	CG2	ILE	A	189	10.928	16.708	30.958	1.00	22.51	C
ATOM	1409	CD1	ILE	A	189	12.918	14.425	31.075	1.00	23.06	C
ATOM	1410	N	GLY	A	190	11.276	20.186	30.673	1.00	22.37	N
ATOM	1411	CA	GLY	A	190	10.587	21.336	31.206	1.00	22.29	C
ATOM	1412	C	GLY	A	190	9.124	21.342	30.902	1.00	22.48	C
ATOM	1413	O	GLY	A	190	8.652	20.633	30.007	1.00	22.39	O
ATOM	1414	N	MET	A	191	8.402	22.134	31.687	1.00	22.74	N

ATOM	1415	CA	MET	A	191	6.991	22.339	31.467	1.00	23.16	C
ATOM	1416	C	MET	A	191	6.838	23.482	30.478	1.00	22.94	C
ATOM	1417	O	MET	A	191	7.738	24.329	30.338	1.00	23.03	O
ATOM	1418	CB	MET	A	191	6.283	22.673	32.784	1.00	23.69	C
ATOM	1419	CG	MET	A	191	6.224	21.513	33.741	1.00	25.02	C
ATOM	1420	SD	MET	A	191	5.664	21.927	35.415	1.00	28.11	S
ATOM	1421	CE	MET	A	191	4.016	22.460	35.097	1.00	28.35	C
ATOM	1422	N	GLU	A	192	5.712	23.492	29.773	1.00	22.60	N
ATOM	1423	CA	GLU	A	192	5.410	24.544	28.810	1.00	22.83	C
ATOM	1424	C	GLU	A	192	5.495	25.895	29.490	1.00	22.51	C
ATOM	1425	O	GLU	A	192	5.062	26.046	30.614	1.00	22.41	O
ATOM	1426	CB	GLU	A	192	4.005	24.342	28.249	1.00	22.86	C
ATOM	1427	CG	GLU	A	192	2.925	24.367	29.315	1.00	24.36	C
ATOM	1428	CD	GLU	A	192	1.572	23.891	28.814	1.00	25.71	C
ATOM	1429	OE1	GLU	A	192	1.503	23.294	27.718	1.00	25.51	O
ATOM	1430	OE2	GLU	A	192	0.582	24.128	29.525	1.00	26.30	O
ATOM	1431	N	GLY	A	193	6.069	26.888	28.828	1.00	22.95	N
ATOM	1432	CA	GLY	A	193	6.185	28.199	29.444	1.00	22.50	C
ATOM	1433	C	GLY	A	193	7.465	28.414	30.254	1.00	22.36	C
ATOM	1434	O	GLY	A	193	7.756	29.544	30.602	1.00	23.08	O
ATOM	1435	N	ASN	A	194	8.219	27.361	30.566	1.00	21.60	N
ATOM	1436	CA	ASN	A	194	9.456	27.506	31.341	1.00	21.44	C
ATOM	1437	C	ASN	A	194	10.489	28.320	30.581	1.00	21.19	C
ATOM	1438	O	ASN	A	194	10.635	28.134	29.372	1.00	22.08	O
ATOM	1439	CB	ASN	A	194	10.099	26.147	31.629	1.00	21.05	C
ATOM	1440	CG	ASN	A	194	9.494	25.435	32.801	1.00	20.86	C
ATOM	1441	OD1	ASN	A	194	8.509	25.883	33.385	1.00	22.48	O
ATOM	1442	ND2	ASN	A	194	10.092	24.312	33.167	1.00	17.74	N
ATOM	1443	N	VAL	A	195	11.213	29.183	31.290	1.00	20.69	N
ATOM	1444	CA	VAL	A	195	12.268	29.993	30.701	1.00	20.84	C
ATOM	1445	C	VAL	A	195	13.572	29.854	31.450	1.00	20.06	C
ATOM	1446	O	VAL	A	195	13.601	29.874	32.681	1.00	20.75	O
ATOM	1447	CB	VAL	A	195	11.903	31.502	30.686	1.00	21.41	C
ATOM	1448	CG1	VAL	A	195	13.081	32.357	30.219	1.00	22.24	C
ATOM	1449	CG2	VAL	A	195	10.666	31.767	29.843	1.00	22.30	C
ATOM	1450	N	THR	A	196	14.651	29.673	30.702	1.00	19.51	N
ATOM	1451	CA	THR	A	196	15.993	29.737	31.257	1.00	19.52	C
ATOM	1452	C	THR	A	196	16.487	31.117	30.820	1.00	19.98	C
ATOM	1453	O	THR	A	196	16.653	31.352	29.620	1.00	19.56	O
ATOM	1454	CB	THR	A	196	16.896	28.677	30.675	1.00	19.26	C
ATOM	1455	OG1	THR	A	196	16.526	27.366	31.162	1.00	21.13	O
ATOM	1456	CG2	THR	A	196	18.309	28.886	31.185	1.00	19.85	C
ATOM	1457	N	PRO	A	197	16.627	32.043	31.767	1.00	20.27	N
ATOM	1458	CA	PRO	A	197	17.046	33.412	31.463	1.00	20.52	C
ATOM	1459	C	PRO	A	197	18.431	33.497	30.836	1.00	20.89	C
ATOM	1460	O	PRO	A	197	19.277	32.609	31.025	1.00	20.66	O
ATOM	1461	CB	PRO	A	197	17.018	34.099	32.825	1.00	21.22	C
ATOM	1462	CG	PRO	A	197	16.144	33.263	33.657	1.00	21.01	C
ATOM	1463	CD	PRO	A	197	16.309	31.872	33.189	1.00	20.28	C
ATOM	1464	N	ALA	A	198	18.633	34.577	30.089	1.00	20.28	N
ATOM	1465	CA	ALA	A	198	19.841	34.817	29.341	1.00	20.22	C
ATOM	1466	C	ALA	A	198	21.130	34.634	30.146	1.00	20.58	C
ATOM	1467	O	ALA	A	198	21.284	35.186	31.235	1.00	19.91	O
ATOM	1468	CB	ALA	A	198	19.791	36.222	28.759	1.00	20.52	C
ATOM	1469	N	HIS	A	199	22.062	33.891	29.563	1.00	20.87	N
ATOM	1470	CA	HIS	A	199	23.371	33.646	30.158	1.00	21.41	C
ATOM	1471	C	HIS	A	199	24.281	33.168	29.063	1.00	21.82	C
ATOM	1472	O	HIS	A	199	23.826	32.892	27.943	1.00	21.80	O
ATOM	1473	CB	HIS	A	199	23.305	32.534	31.198	1.00	21.44	C
ATOM	1474	CG	HIS	A	199	22.915	31.220	30.617	1.00	22.21	C
ATOM	1475	ND1	HIS	A	199	21.619	30.940	30.253	1.00	21.93	N

ATOM	1476	CD2	HIS	A	199	23.644	30.120	30.296	1.00	22.04	C
ATOM	1477	CE1	HIS	A	199	21.561	29.721	29.749	1.00	23.76	C
ATOM	1478	NE2	HIS	A	199	22.777	29.214	29.737	1.00	22.57	N
ATOM	1479	N	TYR	A	200	25.568	33.060	29.384	1.00	22.40	N
ATOM	1480	CA	TYR	A	200	26.536	32.456	28.469	1.00	22.34	C
ATOM	1481	C	TYR	A	200	27.225	31.316	29.197	1.00	22.03	C
ATOM	1482	O	TYR	A	200	27.328	31.330	30.425	1.00	21.90	O
ATOM	1483	CB	TYR	A	200	27.544	33.458	27.924	1.00	22.22	C
ATOM	1484	CG	TYR	A	200	28.517	34.051	28.924	1.00	22.16	C
ATOM	1485	CD1	TYR	A	200	29.746	33.454	29.171	1.00	21.36	C
ATOM	1486	CD2	TYR	A	200	28.236	35.246	29.565	1.00	22.58	C
ATOM	1487	CE1	TYR	A	200	30.638	33.997	30.052	1.00	21.48	C
ATOM	1488	CE2	TYR	A	200	29.128	35.806	30.462	1.00	21.46	C
ATOM	1489	CZ	TYR	A	200	30.334	35.189	30.695	1.00	22.03	C
ATOM	1490	OH	TYR	A	200	31.230	35.733	31.593	1.00	20.69	O
ATOM	1491	N	ASP	A	201	27.681	30.310	28.444	1.00	22.42	N
ATOM	1492	CA	ASP	A	201	28.381	29.152	29.048	1.00	22.10	C
ATOM	1493	C	ASP	A	201	29.801	29.142	28.531	1.00	21.96	C
ATOM	1494	O	ASP	A	201	30.018	29.601	27.433	1.00	22.26	O
ATOM	1495	CB	ASP	A	201	27.722	27.840	28.661	1.00	21.97	C
ATOM	1496	CG	ASP	A	201	26.311	27.714	29.181	1.00	21.99	C
ATOM	1497	OD1	ASP	A	201	26.126	27.649	30.421	1.00	22.61	O
ATOM	1498	OD2	ASP	A	201	25.330	27.624	28.412	1.00	19.35	O
ATOM	1499	N	GLU	A	202	30.769	28.620	29.283	1.00	21.66	N
ATOM	1500	CA	GLU	A	202	32.146	28.612	28.773	1.00	23.16	C
ATOM	1501	C	GLU	A	202	32.515	27.334	28.035	1.00	23.38	C
ATOM	1502	O	GLU	A	202	33.684	26.951	28.049	1.00	25.48	O
ATOM	1503	CB	GLU	A	202	33.178	28.774	29.897	1.00	23.46	C
ATOM	1504	CG	GLU	A	202	33.110	30.064	30.681	1.00	26.52	C
ATOM	1505	CD	GLU	A	202	34.233	30.132	31.704	1.00	28.11	C
ATOM	1506	OE1	GLU	A	202	34.207	29.387	32.698	1.00	29.06	O
ATOM	1507	OE2	GLU	A	202	35.148	30.920	31.490	1.00	31.04	O
ATOM	1508	N	GLN	A	203	31.538	26.632	27.476	1.00	22.70	N
ATOM	1509	CA	GLN	A	203	31.808	25.451	26.685	1.00	22.01	C
ATOM	1510	C	GLN	A	203	31.049	25.576	25.373	1.00	21.93	C
ATOM	1511	O	GLN	A	203	30.113	26.378	25.253	1.00	22.49	O
ATOM	1512	CB	GLN	A	203	31.409	24.169	27.427	1.00	22.02	C
ATOM	1513	CG	GLN	A	203	32.287	23.822	28.657	1.00	22.33	C
ATOM	1514	CD	GLN	A	203	32.103	22.379	29.165	1.00	23.14	C
ATOM	1515	OE1	GLN	A	203	31.948	21.448	28.370	1.00	21.46	O
ATOM	1516	NE2	GLN	A	203	32.138	22.203	30.489	1.00	21.21	N
ATOM	1517	N	GLN	A	204	31.486	24.807	24.380	1.00	21.81	N
ATOM	1518	CA	GLN	A	204	30.806	24.720	23.092	1.00	21.45	C
ATOM	1519	C	GLN	A	204	29.610	23.799	23.295	1.00	21.12	C
ATOM	1520	O	GLN	A	204	29.673	22.862	24.088	1.00	20.29	O
ATOM	1521	CB	GLN	A	204	31.740	24.140	22.040	1.00	21.75	C
ATOM	1522	CG	GLN	A	204	33.033	24.928	21.852	1.00	21.56	C
ATOM	1523	CD	GLN	A	204	32.856	26.198	21.000	1.00	21.57	C
ATOM	1524	OE1	GLN	A	204	31.741	26.609	20.692	1.00	20.30	O
ATOM	1525	NE2	GLN	A	204	33.966	26.776	20.594	1.00	18.60	N
ATOM	1526	N	ASN	A	205	28.522	24.060	22.580	1.00	21.25	N
ATOM	1527	CA	ASN	A	205	27.291	23.316	22.795	1.00	21.04	C
ATOM	1528	C	ASN	A	205	26.529	22.977	21.526	1.00	21.25	C
ATOM	1529	O	ASN	A	205	26.160	23.880	20.747	1.00	20.22	O
ATOM	1530	CB	ASN	A	205	26.387	24.194	23.654	1.00	21.95	C
ATOM	1531	CG	ASN	A	205	25.116	23.507	24.098	1.00	21.98	C
ATOM	1532	OD1	ASN	A	205	24.759	22.418	23.643	1.00	21.21	O
ATOM	1533	ND2	ASN	A	205	24.400	24.178	24.979	1.00	17.72	N
ATOM	1534	N	PHE	A	206	26.314	21.677	21.319	1.00	19.90	N
ATOM	1535	CA	PHE	A	206	25.339	21.230	20.354	1.00	19.77	C
ATOM	1536	C	PHE	A	206	24.085	20.869	21.150	1.00	19.25	C

ATOM	1537	O	PHE	A	206	24.094	19.939	21.959	1.00	19.15	O
ATOM	1538	CB	PHE	A	206	25.849	20.049	19.545	1.00	20.09	C
ATOM	1539	CG	PHE	A	206	26.786	20.442	18.454	1.00	19.77	C
ATOM	1540	CD1	PHE	A	206	26.371	21.294	17.453	1.00	22.21	C
ATOM	1541	CD2	PHE	A	206	28.079	19.973	18.435	1.00	20.21	C
ATOM	1542	CE1	PHE	A	206	27.213	21.643	16.434	1.00	22.06	C
ATOM	1543	CE2	PHE	A	206	28.941	20.336	17.418	1.00	20.66	C
ATOM	1544	CZ	PHE	A	206	28.504	21.190	16.427	1.00	22.18	C
ATOM	1545	N	PHE	A	207	23.009	21.596	20.869	1.00	18.42	N
ATOM	1546	CA	PHE	A	207	21.760	21.590	21.620	1.00	18.78	C
ATOM	1547	C	PHE	A	207	20.732	20.841	20.804	1.00	19.36	C
ATOM	1548	O	PHE	A	207	20.241	21.359	19.818	1.00	19.26	O
ATOM	1549	CB	PHE	A	207	21.366	23.063	21.836	1.00	18.70	C
ATOM	1550	CG	PHE	A	207	20.120	23.301	22.635	1.00	17.94	C
ATOM	1551	CD1	PHE	A	207	18.899	23.404	22.007	1.00	17.78	C
ATOM	1552	CD2	PHE	A	207	20.188	23.548	23.989	1.00	18.50	C
ATOM	1553	CE1	PHE	A	207	17.772	23.684	22.713	1.00	19.52	C
ATOM	1554	CE2	PHE	A	207	19.052	23.823	24.705	1.00	20.66	C
ATOM	1555	CZ	PHE	A	207	17.836	23.909	24.053	1.00	19.44	C
ATOM	1556	N	ALA	A	208	20.435	19.618	21.234	1.00	19.90	N
ATOM	1557	CA	ALA	A	208	19.628	18.670	20.469	1.00	20.17	C
ATOM	1558	C	ALA	A	208	18.210	18.541	20.991	1.00	20.52	C
ATOM	1559	O	ALA	A	208	17.971	17.946	22.047	1.00	20.61	O
ATOM	1560	CB	ALA	A	208	20.294	17.308	20.494	1.00	19.33	C
ATOM	1561	N	GLN	A	209	17.270	19.076	20.219	1.00	20.94	N
ATOM	1562	CA	GLN	A	209	15.880	19.062	20.627	1.00	20.82	C
ATOM	1563	C	GLN	A	209	15.227	17.712	20.323	1.00	20.75	C
ATOM	1564	O	GLN	A	209	15.401	17.129	19.224	1.00	19.59	O
ATOM	1565	CB	GLN	A	209	15.141	20.216	19.961	1.00	20.91	C
ATOM	1566	CG	GLN	A	209	13.735	20.431	20.463	1.00	20.52	C
ATOM	1567	CD	GLN	A	209	13.673	20.840	21.928	1.00	20.69	C
ATOM	1568	OE1	GLN	A	209	14.702	21.155	22.562	1.00	19.72	O
ATOM	1569	NE2	GLN	A	209	12.460	20.852	22.473	1.00	19.21	N
ATOM	1570	N	ILE	A	210	14.442	17.248	21.296	1.00	20.72	N
ATOM	1571	CA	ILE	A	210	13.853	15.922	21.252	1.00	21.43	C
ATOM	1572	C	ILE	A	210	12.334	15.911	21.292	1.00	21.72	C
ATOM	1573	O	ILE	A	210	11.728	15.195	20.534	1.00	22.12	O
ATOM	1574	CB	ILE	A	210	14.396	15.095	22.424	1.00	21.84	C
ATOM	1575	CG1	ILE	A	210	15.859	14.733	22.156	1.00	22.48	C
ATOM	1576	CG2	ILE	A	210	13.581	13.832	22.622	1.00	21.83	C
ATOM	1577	CD1	ILE	A	210	16.631	14.354	23.382	1.00	24.34	C
ATOM	1578	N	LYS	A	211	11.728	16.677	22.188	1.00	22.08	N
ATOM	1579	CA	LYS	A	211	10.279	16.715	22.319	1.00	21.72	C
ATOM	1580	C	LYS	A	211	9.858	18.130	22.564	1.00	21.67	C
ATOM	1581	O	LYS	A	211	10.468	18.841	23.372	1.00	20.87	O
ATOM	1582	CB	LYS	A	211	9.797	15.856	23.487	1.00	22.44	C
ATOM	1583	CG	LYS	A	211	8.267	15.562	23.479	1.00	23.29	C
ATOM	1584	CD	LYS	A	211	7.791	15.022	24.824	1.00	25.02	C
ATOM	1585	CE	LYS	A	211	6.494	14.215	24.757	1.00	26.00	C
ATOM	1586	NZ	LYS	A	211	5.561	14.522	23.634	1.00	26.43	N
ATOM	1587	N	GLY	A	212	8.798	18.541	21.875	1.00	21.35	N
ATOM	1588	CA	GLY	A	212	8.306	19.891	21.994	1.00	21.71	C
ATOM	1589	C	GLY	A	212	9.195	20.906	21.297	1.00	21.63	C
ATOM	1590	O	GLY	A	212	10.150	20.572	20.591	1.00	21.32	O
ATOM	1591	N	TYR	A	213	8.871	22.166	21.522	1.00	22.35	N
ATOM	1592	CA	TYR	A	213	9.533	23.279	20.862	1.00	22.42	C
ATOM	1593	C	TYR	A	213	10.028	24.293	21.868	1.00	22.33	C
ATOM	1594	O	TYR	A	213	9.340	24.589	22.853	1.00	21.55	O
ATOM	1595	CB	TYR	A	213	8.556	23.951	19.916	1.00	23.46	C
ATOM	1596	CG	TYR	A	213	8.114	23.034	18.815	1.00	24.81	C
ATOM	1597	CD1	TYR	A	213	7.100	22.112	19.020	1.00	28.16	C

ATOM	1598	CD2	TYR	A	213	8.751	23.051	17.589	1.00	26.74	C
ATOM	1599	CE1	TYR	A	213	6.716	21.237	18.011	1.00	28.91	C
ATOM	1600	CE2	TYR	A	213	8.378	22.193	16.585	1.00	28.63	C
ATOM	1601	CZ	TYR	A	213	7.366	21.295	16.795	1.00	29.44	C
ATOM	1602	OH	TYR	A	213	7.013	20.456	15.756	1.00	33.67	O
ATOM	1603	N	LYS	A	214	11.239	24.788	21.609	1.00	21.60	N
ATOM	1604	CA	LYS	A	214	11.875	25.793	22.414	1.00	21.89	C
ATOM	1605	C	LYS	A	214	12.312	26.947	21.528	1.00	21.62	C
ATOM	1606	O	LYS	A	214	12.878	26.747	20.442	1.00	22.07	O
ATOM	1607	CB	LYS	A	214	13.103	25.239	23.140	1.00	22.10	C
ATOM	1608	CG	LYS	A	214	12.796	24.385	24.338	1.00	22.63	C
ATOM	1609	CD	LYS	A	214	14.103	23.893	24.964	1.00	24.24	C
ATOM	1610	CE	LYS	A	214	13.923	23.478	26.402	1.00	23.54	C
ATOM	1611	NZ	LYS	A	214	15.169	22.872	26.951	1.00	22.10	N
ATOM	1612	N	ARG	A	215	12.007	28.156	21.974	1.00	20.97	N
ATOM	1613	CA	ARG	A	215	12.447	29.347	21.277	1.00	20.73	C
ATOM	1614	C	ARG	A	215	13.778	29.724	21.900	1.00	20.03	C
ATOM	1615	O	ARG	A	215	13.885	29.828	23.113	1.00	19.70	O
ATOM	1616	CB	ARG	A	215	11.461	30.470	21.493	1.00	21.30	C
ATOM	1617	CG	ARG	A	215	11.754	31.760	20.726	1.00	22.21	C
ATOM	1618	CD	ARG	A	215	11.177	32.925	21.473	1.00	24.71	C
ATOM	1619	NE	ARG	A	215	11.122	34.156	20.715	1.00	25.47	N
ATOM	1620	CZ	ARG	A	215	10.479	35.235	21.130	1.00	26.28	C
ATOM	1621	NH1	ARG	A	215	9.844	35.245	22.299	1.00	26.07	N
ATOM	1622	NH2	ARG	A	215	10.483	36.314	20.384	1.00	28.83	N
ATOM	1623	N	CYS	A	216	14.794	29.877	21.074	1.00	19.45	N
ATOM	1624	CA	CYS	A	216	16.116	30.219	21.554	1.00	19.58	C
ATOM	1625	C	CYS	A	216	16.503	31.587	20.993	1.00	19.92	C
ATOM	1626	O	CYS	A	216	16.503	31.774	19.789	1.00	20.16	O
ATOM	1627	CB	CYS	A	216	17.110	29.186	21.073	1.00	20.01	C
ATOM	1628	SG	CYS	A	216	16.693	27.450	21.462	1.00	21.39	S
ATOM	1629	N	ILE	A	217	16.812	32.530	21.878	1.00	19.28	N
ATOM	1630	CA	ILE	A	217	17.286	33.845	21.504	1.00	19.58	C
ATOM	1631	C	ILE	A	217	18.742	33.961	21.937	1.00	19.21	C
ATOM	1632	O	ILE	A	217	19.055	33.849	23.126	1.00	19.05	O
ATOM	1633	CB	ILE	A	217	16.424	34.943	22.168	1.00	19.88	C
ATOM	1634	CG1	ILE	A	217	14.926	34.687	21.878	1.00	20.56	C
ATOM	1635	CG2	ILE	A	217	16.803	36.312	21.627	1.00	20.25	C
ATOM	1636	CD1	ILE	A	217	13.984	35.692	22.488	1.00	20.99	C
ATOM	1637	N	LEU	A	218	19.620	34.176	20.965	1.00	19.01	N
ATOM	1638	CA	LEU	A	218	21.048	34.271	21.222	1.00	19.61	C
ATOM	1639	C	LEU	A	218	21.568	35.682	20.970	1.00	20.43	C
ATOM	1640	O	LEU	A	218	21.018	36.414	20.136	1.00	19.76	O
ATOM	1641	CB	LEU	A	218	21.816	33.329	20.308	1.00	19.66	C
ATOM	1642	CG	LEU	A	218	21.906	31.881	20.778	1.00	19.74	C
ATOM	1643	CD1	LEU	A	218	20.549	31.237	20.905	1.00	20.56	C
ATOM	1644	CD2	LEU	A	218	22.733	31.099	19.798	1.00	21.32	C
ATOM	1645	N	PHE	A	219	22.626	36.046	21.700	1.00	20.11	N
ATOM	1646	CA	PHE	A	219	23.282	37.324	21.524	1.00	20.96	C
ATOM	1647	C	PHE	A	219	24.805	37.078	21.453	1.00	21.19	C
ATOM	1648	O	PHE	A	219	25.351	36.313	22.257	1.00	21.20	O
ATOM	1649	CB	PHE	A	219	22.942	38.304	22.678	1.00	20.97	C
ATOM	1650	CG	PHE	A	219	21.463	38.456	22.954	1.00	20.44	C
ATOM	1651	CD1	PHE	A	219	20.821	37.637	23.846	1.00	21.46	C
ATOM	1652	CD2	PHE	A	219	20.724	39.447	22.334	1.00	22.49	C
ATOM	1653	CE1	PHE	A	219	19.472	37.790	24.108	1.00	22.33	C
ATOM	1654	CE2	PHE	A	219	19.387	39.605	22.605	1.00	21.32	C
ATOM	1655	CZ	PHE	A	219	18.760	38.773	23.480	1.00	21.50	C
ATOM	1656	N	PRO	A	220	25.485	37.705	20.498	1.00	21.20	N
ATOM	1657	CA	PRO	A	220	26.936	37.541	20.368	1.00	21.72	C
ATOM	1658	C	PRO	A	220	27.683	38.108	21.569	1.00	21.31	C

ATOM	1659	O	PRO	A	220	27.165	38.972	22.280	1.00	20.90	O
ATOM	1660	CB	PRO	A	220	27.302	38.341	19.120	1.00	21.41	C
ATOM	1661	CG	PRO	A	220	26.020	38.781	18.510	1.00	23.12	C
ATOM	1662	CD	PRO	A	220	24.938	38.637	19.508	1.00	22.22	C
ATOM	1663	N	PRO	A	221	28.884	37.594	21.806	1.00	21.46	N
ATOM	1664	CA	PRO	A	221	29.721	38.053	22.918	1.00	21.33	C
ATOM	1665	C	PRO	A	221	29.961	39.556	22.940	1.00	21.04	C
ATOM	1666	O	PRO	A	221	30.220	40.096	24.009	1.00	20.68	O
ATOM	1667	CB	PRO	A	221	31.032	37.325	22.681	1.00	21.42	C
ATOM	1668	CG	PRO	A	221	30.656	36.134	21.939	1.00	22.42	C
ATOM	1669	CD	PRO	A	221	29.522	36.510	21.041	1.00	21.53	C
ATOM	1670	N	ASP	A	222	29.877	40.225	21.792	1.00	21.47	N
ATOM	1671	CA	ASP	A	222	30.128	41.669	21.749	1.00	21.80	C
ATOM	1672	C	ASP	A	222	28.946	42.464	22.271	1.00	21.74	C
ATOM	1673	O	ASP	A	222	28.970	43.685	22.261	1.00	22.16	O
ATOM	1674	CB	ASP	A	222	30.568	42.148	20.355	1.00	21.85	C
ATOM	1675	CG	ASP	A	222	29.433	42.151	19.325	1.00	24.83	C
ATOM	1676	OD1	ASP	A	222	28.311	41.668	19.585	1.00	25.32	O
ATOM	1677	OD2	ASP	A	222	29.595	42.607	18.186	1.00	29.30	O
ATOM	1678	N	GLN	A	223	27.916	41.775	22.748	1.00	21.94	N
ATOM	1679	CA	GLN	A	223	26.794	42.453	23.388	1.00	22.17	C
ATOM	1680	C	GLN	A	223	26.897	42.388	24.928	1.00	21.93	C
ATOM	1681	O	GLN	A	223	25.926	42.681	25.660	1.00	21.70	O
ATOM	1682	CB	GLN	A	223	25.456	41.911	22.832	1.00	22.84	C
ATOM	1683	CG	GLN	A	223	25.149	42.508	21.412	1.00	25.34	C
ATOM	1684	CD	GLN	A	223	23.728	42.283	20.965	1.00	27.81	C
ATOM	1685	OE1	GLN	A	223	22.801	42.473	21.750	1.00	30.00	O
ATOM	1686	NE2	GLN	A	223	23.543	41.850	19.709	1.00	29.69	N
ATOM	1687	N	PHE	A	224	28.079	42.027	25.417	1.00	21.04	N
ATOM	1688	CA	PHE	A	224	28.363	42.035	26.860	1.00	21.41	C
ATOM	1689	C	PHE	A	224	27.886	43.350	27.527	1.00	21.57	C
ATOM	1690	O	PHE	A	224	27.240	43.315	28.561	1.00	21.34	O
ATOM	1691	CB	PHE	A	224	29.883	41.874	27.076	1.00	20.95	C
ATOM	1692	CG	PHE	A	224	30.304	41.773	28.529	1.00	21.82	C
ATOM	1693	CD1	PHE	A	224	30.452	42.912	29.319	1.00	19.33	C
ATOM	1694	CD2	PHE	A	224	30.595	40.552	29.089	1.00	20.01	C
ATOM	1695	CE1	PHE	A	224	30.870	42.807	30.630	1.00	20.89	C
ATOM	1696	CE2	PHE	A	224	31.025	40.449	30.397	1.00	20.59	C
ATOM	1697	CZ	PHE	A	224	31.151	41.579	31.171	1.00	20.99	C
ATOM	1698	N	GLU	A	225	28.195	44.495	26.916	1.00	22.10	N
ATOM	1699	CA	GLU	A	225	27.823	45.800	27.466	1.00	23.01	C
ATOM	1700	C	GLU	A	225	26.337	46.042	27.574	1.00	22.26	C
ATOM	1701	O	GLU	A	225	25.928	46.946	28.311	1.00	21.91	O
ATOM	1702	CB	GLU	A	225	28.401	46.935	26.624	1.00	24.27	C
ATOM	1703	CG	GLU	A	225	29.886	47.133	26.857	1.00	29.26	C
ATOM	1704	CD	GLU	A	225	30.312	48.579	27.062	1.00	35.91	C
ATOM	1705	OE1	GLU	A	225	29.745	49.317	27.942	1.00	39.92	O
ATOM	1706	OE2	GLU	A	225	31.281	48.958	26.368	1.00	40.78	O
ATOM	1707	N	CYS	A	226	25.539	45.258	26.836	1.00	21.15	N
ATOM	1708	CA	CYS	A	226	24.081	45.396	26.851	1.00	21.01	C
ATOM	1709	C	CYS	A	226	23.359	44.417	27.768	1.00	20.67	C
ATOM	1710	O	CYS	A	226	22.159	44.601	28.043	1.00	18.85	O
ATOM	1711	CB	CYS	A	226	23.520	45.190	25.447	1.00	20.88	C
ATOM	1712	SG	CYS	A	226	24.194	46.296	24.190	1.00	22.39	S
ATOM	1713	N	LEU	A	227	24.085	43.399	28.256	1.00	20.56	N
ATOM	1714	CA	LEU	A	227	23.461	42.304	28.985	1.00	20.48	C
ATOM	1715	C	LEU	A	227	23.795	42.185	30.466	1.00	20.42	C
ATOM	1716	O	LEU	A	227	23.180	41.398	31.187	1.00	20.78	O
ATOM	1717	CB	LEU	A	227	23.737	41.003	28.232	1.00	21.20	C
ATOM	1718	CG	LEU	A	227	22.865	40.993	26.963	1.00	22.96	C
ATOM	1719	CD1	LEU	A	227	23.373	40.023	25.929	1.00	24.99	C

ATOM	1720	CD2	LEU	A	227	21.420	40.655	27.323	1.00	24.79
ATOM	1721	N	TYR	A	228	24.793	42.939	30.901	1.00	20.08
ATOM	1722	CA	TYR	A	228	25.060	43.127	32.303	1.00	19.66
ATOM	1723	C	TYR	A	228	25.061	41.858	33.165	1.00	19.95
ATOM	1724	O	TYR	A	228	24.229	41.697	34.058	1.00	19.69
ATOM	1725	CB	TYR	A	228	24.050	44.137	32.857	1.00	19.77
ATOM	1726	CG	TYR	A	228	24.028	45.481	32.119	1.00	19.37
ATOM	1727	CD1	TYR	A	228	24.836	46.513	32.515	1.00	18.65
ATOM	1728	CD2	TYR	A	228	23.180	45.697	31.036	1.00	19.82
ATOM	1729	CE1	TYR	A	228	24.819	47.759	31.862	1.00	20.56
ATOM	1730	CE2	TYR	A	228	23.143	46.920	30.380	1.00	20.78
ATOM	1731	CZ	TYR	A	228	23.962	47.956	30.801	1.00	20.88
ATOM	1732	OH	TYR	A	228	23.944	49.174	30.152	1.00	19.73
ATOM	1733	N	PRO	A	229	26.028	40.981	32.941	1.00	19.99
ATOM	1734	CA	PRO	A	229	26.140	39.781	33.768	1.00	20.44
ATOM	1735	C	PRO	A	229	26.481	40.141	35.196	1.00	19.99
ATOM	1736	O	PRO	A	229	27.130	41.166	35.451	1.00	20.03
ATOM	1737	CB	PRO	A	229	27.334	39.043	33.157	1.00	20.11
ATOM	1738	CG	PRO	A	229	28.125	40.130	32.541	1.00	21.04
ATOM	1739	CD	PRO	A	229	27.099	41.050	31.935	1.00	19.79
ATOM	1740	N	TYR	A	230	26.044	39.294	36.115	1.00	19.50
ATOM	1741	CA	TYR	A	230	26.348	39.458	37.511	1.00	19.17
ATOM	1742	C	TYR	A	230	27.860	39.464	37.704	1.00	19.14
ATOM	1743	O	TYR	A	230	28.598	39.060	36.832	1.00	18.83
ATOM	1744	CB	TYR	A	230	25.746	38.313	38.320	1.00	18.51
ATOM	1745	CG	TYR	A	230	24.253	38.411	38.531	1.00	18.95
ATOM	1746	CD1	TYR	A	230	23.368	37.949	37.560	1.00	18.39
ATOM	1747	CD2	TYR	A	230	23.719	38.969	39.707	1.00	18.65
ATOM	1748	CE1	TYR	A	230	22.001	38.006	37.751	1.00	17.34
ATOM	1749	CE2	TYR	A	230	22.337	39.043	39.903	1.00	17.05
ATOM	1750	CZ	TYR	A	230	21.492	38.550	38.909	1.00	17.79
ATOM	1751	OH	TYR	A	230	20.127	38.601	39.051	1.00	19.77
ATOM	1752	N	PRO	A	231	28.316	39.963	38.843	1.00	19.95
ATOM	1753	CA	PRO	A	231	29.722	39.829	39.213	1.00	20.42
ATOM	1754	C	PRO	A	231	30.157	38.339	39.187	1.00	20.93
ATOM	1755	O	PRO	A	231	29.345	37.463	39.506	1.00	20.71
ATOM	1756	CB	PRO	A	231	29.755	40.369	40.642	1.00	20.70
ATOM	1757	CG	PRO	A	231	28.588	41.341	40.716	1.00	20.76
ATOM	1758	CD	PRO	A	231	27.528	40.701	39.845	1.00	19.53
ATOM	1759	N	VAL	A	232	31.413	38.067	38.831	1.00	20.93
ATOM	1760	CA	VAL	A	232	31.906	36.685	38.742	1.00	21.31
ATOM	1761	C	VAL	A	232	31.722	35.843	40.014	1.00	20.95
ATOM	1762	O	VAL	A	232	31.431	34.650	39.934	1.00	20.50
ATOM	1763	CB	VAL	A	232	33.398	36.655	38.304	1.00	21.35
ATOM	1764	CG1	VAL	A	232	34.029	35.280	38.552	1.00	22.23
ATOM	1765	CG2	VAL	A	232	33.533	37.031	36.850	1.00	21.71
ATOM	1766	N	HIS	A	233	31.872	36.455	41.181	1.00	21.10
ATOM	1767	CA	HIS	A	233	31.742	35.721	42.438	1.00	21.01
ATOM	1768	C	HIS	A	233	30.302	35.571	42.944	1.00	20.95
ATOM	1769	O	HIS	A	233	30.045	34.864	43.914	1.00	19.92
ATOM	1770	CB	HIS	A	233	32.606	36.363	43.501	1.00	21.12
ATOM	1771	CG	HIS	A	233	34.063	36.342	43.174	1.00	21.52
ATOM	1772	ND1	HIS	A	233	34.720	37.424	42.625	1.00	22.78
ATOM	1773	CD2	HIS	A	233	34.993	35.368	43.318	1.00	22.27
ATOM	1774	CE1	HIS	A	233	35.994	37.114	42.446	1.00	24.00
ATOM	1775	NE2	HIS	A	233	36.186	35.870	42.857	1.00	23.34
ATOM	1776	N	HIS	A	234	29.364	36.236	42.288	1.00	20.94
ATOM	1777	CA	HIS	A	234	27.959	36.081	42.630	1.00	21.27
ATOM	1778	C	HIS	A	234	27.493	34.722	42.103	1.00	21.28
ATOM	1779	O	HIS	A	234	28.022	34.233	41.107	1.00	21.02
ATOM	1780	CB	HIS	A	234	27.169	37.203	41.989	1.00	21.52

[illegible]

ATOM	1781	CG	HIS	A	234	25.730	37.253	42.374	1.00	21.54	C
ATOM	1782	ND1	HIS	A	234	24.769	36.498	41.740	1.00	21.95	N
ATOM	1783	CD2	HIS	A	234	25.073	38.028	43.269	1.00	22.67	C
ATOM	1784	CE1	HIS	A	234	23.585	36.779	42.254	1.00	22.38	C
ATOM	1785	NE2	HIS	A	234	23.743	37.700	43.187	1.00	21.91	N
ATOM	1786	N	PRO	A	235	26.551	34.087	42.792	1.00	21.11	N
ATOM	1787	CA	PRO	A	235	25.999	32.809	42.338	1.00	21.46	C
ATOM	1788	C	PRO	A	235	25.461	32.835	40.916	1.00	21.10	C
ATOM	1789	O	PRO	A	235	25.517	31.802	40.244	1.00	20.37	O
ATOM	1790	CB	PRO	A	235	24.860	32.551	43.325	1.00	21.90	C
ATOM	1791	CG	PRO	A	235	25.316	33.240	44.574	1.00	22.00	C
ATOM	1792	CD	PRO	A	235	26.007	34.485	44.100	1.00	21.41	C
ATOM	1793	N	CYS	A	236	24.997	33.989	40.452	1.00	20.41	N
ATOM	1794	CA	CYS	A	236	24.456	34.077	39.110	1.00	20.17	C
ATOM	1795	C	CYS	A	236	25.503	34.554	38.107	1.00	20.00	C
ATOM	1796	O	CYS	A	236	25.180	35.111	37.061	1.00	20.30	O
ATOM	1797	CB	CYS	A	236	23.195	34.924	39.090	1.00	19.66	C
ATOM	1798	SG	CYS	A	236	21.914	34.209	40.133	1.00	20.71	S
ATOM	1799	N	ASP	A	237	26.768	34.311	38.433	1.00	19.87	N
ATOM	1800	CA	ASP	A	237	27.857	34.498	37.485	1.00	19.60	C
ATOM	1801	C	ASP	A	237	27.459	33.915	36.105	1.00	19.48	C
ATOM	1802	O	ASP	A	237	26.883	32.834	36.020	1.00	18.94	O
ATOM	1803	CB	ASP	A	237	29.075	33.789	38.031	1.00	19.20	C
ATOM	1804	CG	ASP	A	237	30.268	33.842	37.107	1.00	20.58	C
ATOM	1805	OD1	ASP	A	237	30.477	34.871	36.373	1.00	18.56	O
ATOM	1806	OD2	ASP	A	237	31.070	32.868	37.077	1.00	19.51	O
ATOM	1807	N	ARG	A	238	27.749	34.675	35.056	1.00	19.85	N
ATOM	1808	CA	ARG	A	238	27.459	34.332	33.664	1.00	20.90	C
ATOM	1809	C	ARG	A	238	26.002	34.572	33.247	1.00	20.67	C
ATOM	1810	O	ARG	A	238	25.709	34.491	32.059	1.00	21.20	O
ATOM	1811	CB	ARG	A	238	27.854	32.876	33.319	1.00	21.44	C
ATOM	1812	CG	ARG	A	238	29.330	32.585	33.482	1.00	21.04	C
ATOM	1813	CD	ARG	A	238	29.710	31.150	33.145	1.00	22.38	C
ATOM	1814	NE	ARG	A	238	29.080	30.243	34.085	1.00	23.86	N
ATOM	1815	CZ	ARG	A	238	27.937	29.584	33.874	1.00	25.81	C
ATOM	1816	NH1	ARG	A	238	27.262	29.686	32.712	1.00	24.18	N
ATOM	1817	NH2	ARG	A	238	27.471	28.810	34.849	1.00	27.18	N
ATOM	1818	N	GLN	A	239	25.105	34.845	34.190	1.00	19.94	N
ATOM	1819	CA	GLN	A	239	23.722	35.141	33.833	1.00	20.58	C
ATOM	1820	C	GLN	A	239	23.535	36.661	33.740	1.00	20.50	C
ATOM	1821	O	GLN	A	239	24.219	37.402	34.446	1.00	20.63	O
ATOM	1822	CB	GLN	A	239	22.735	34.591	34.863	1.00	20.32	C
ATOM	1823	CG	GLN	A	239	23.159	33.314	35.594	1.00	20.84	C
ATOM	1824	CD	GLN	A	239	23.367	32.137	34.673	1.00	21.23	C
ATOM	1825	OE1	GLN	A	239	22.434	31.675	34.014	1.00	21.22	O
ATOM	1826	NE2	GLN	A	239	24.591	31.655	34.616	1.00	20.93	N
ATOM	1827	N	SER	A	240	22.600	37.112	32.900	1.00	20.07	N
ATOM	1828	CA	SER	A	240	22.282	38.533	32.768	1.00	20.18	C
ATOM	1829	C	SER	A	240	21.498	39.027	33.969	1.00	20.71	C
ATOM	1830	O	SER	A	240	20.619	38.316	34.467	1.00	20.85	O
ATOM	1831	CB	SER	A	240	21.405	38.776	31.539	1.00	20.29	C
ATOM	1832	OG	SER	A	240	21.007	40.137	31.454	1.00	21.43	O
ATOM	1833	N	GLN	A	241	21.793	40.246	34.426	1.00	20.46	N
ATOM	1834	CA	GLN	A	241	21.013	40.839	35.496	1.00	20.42	C
ATOM	1835	C	GLN	A	241	19.711	41.433	34.965	1.00	20.26	C
ATOM	1836	O	GLN	A	241	18.839	41.767	35.726	1.00	19.62	O
ATOM	1837	CB	GLN	A	241	21.770	41.949	36.201	1.00	20.32	C
ATOM	1838	CG	GLN	A	241	23.019	41.544	36.912	1.00	20.58	C
ATOM	1839	CD	GLN	A	241	23.771	42.767	37.423	1.00	21.94	C
ATOM	1840	OE1	GLN	A	241	23.524	43.210	38.514	1.00	23.52	O
ATOM	1841	NE2	GLN	A	241	24.670	43.304	36.624	1.00	20.84	N

ATOM	1842	N	VAL	A	242	19.557	41.563	33.659	1.00	20.57
ATOM	1843	CA	VAL	A	242	18.361	42.229	33.182	1.00	20.73
ATOM	1844	C	VAL	A	242	17.173	41.309	33.246	1.00	20.69
ATOM	1845	O	VAL	A	242	17.244	40.190	32.757	1.00	21.13
ATOM	1846	CB	VAL	A	242	18.498	42.667	31.712	1.00	21.03
ATOM	1847	CG1	VAL	A	242	17.204	43.377	31.250	1.00	20.85
ATOM	1848	CG2	VAL	A	242	19.723	43.527	31.498	1.00	20.71
ATOM	1849	N	ASP	A	243	16.071	41.786	33.813	1.00	20.62
ATOM	1850	CA	ASP	A	243	14.807	41.042	33.788	1.00	20.64
ATOM	1851	C	ASP	A	243	14.148	41.348	32.448	1.00	20.94
ATOM	1852	O	ASP	A	243	13.591	42.438	32.230	1.00	19.75
ATOM	1853	CB	ASP	A	243	13.916	41.471	34.963	1.00	20.88
ATOM	1854	CG	ASP	A	243	12.526	40.894	34.898	1.00	21.39
ATOM	1855	OD1	ASP	A	243	12.209	40.153	33.941	1.00	23.25
ATOM	1856	OD2	ASP	A	243	11.672	41.130	35.784	1.00	23.49
ATOM	1857	N	PHE	A	244	14.238	40.385	31.541	1.00	21.19
ATOM	1858	CA	PHE	A	244	13.717	40.549	30.197	1.00	21.56
ATOM	1859	C	PHE	A	244	12.225	40.916	30.204	1.00	22.49
ATOM	1860	O	PHE	A	244	11.734	41.521	29.260	1.00	21.85
ATOM	1861	CB	PHE	A	244	13.940	39.275	29.378	1.00	21.85
ATOM	1862	CG	PHE	A	244	15.316	39.152	28.743	1.00	20.87
ATOM	1863	CD1	PHE	A	244	16.457	39.540	29.399	1.00	20.74
ATOM	1864	CD2	PHE	A	244	15.446	38.617	27.483	1.00	21.00
ATOM	1865	CE1	PHE	A	244	17.691	39.407	28.810	1.00	19.77
ATOM	1866	CE2	PHE	A	244	16.687	38.470	26.890	1.00	20.94
ATOM	1867	CZ	PHE	A	244	17.805	38.881	27.551	1.00	20.34
ATOM	1868	N	ASP	A	245	11.500	40.554	31.259	1.00	23.56
ATOM	1869	CA	ASP	A	245	10.072	40.831	31.311	1.00	24.26
ATOM	1870	C	ASP	A	245	9.794	42.256	31.814	1.00	24.66
ATOM	1871	O	ASP	A	245	8.711	42.762	31.607	1.00	24.21
ATOM	1872	CB	ASP	A	245	9.353	39.841	32.222	1.00	24.74
ATOM	1873	CG	ASP	A	245	9.408	38.427	31.721	1.00	26.56
ATOM	1874	OD1	ASP	A	245	9.368	38.210	30.493	1.00	28.81
ATOM	1875	OD2	ASP	A	245	9.489	37.455	32.498	1.00	28.39
ATOM	1876	N	ASN	A	246	10.758	42.876	32.492	1.00	24.80
ATOM	1877	CA	ASN	A	246	10.605	44.244	32.989	1.00	25.56
ATOM	1878	C	ASN	A	246	11.960	44.902	33.106	1.00	25.31
ATOM	1879	O	ASN	A	246	12.500	45.064	34.205	1.00	25.43
ATOM	1880	CB	ASN	A	246	9.912	44.285	34.345	1.00	26.09
ATOM	1881	CG	ASN	A	246	9.555	45.716	34.765	1.00	29.59
ATOM	1882	OD1	ASN	A	246	9.366	46.598	33.912	1.00	32.67
ATOM	1883	ND2	ASN	A	246	9.468	45.954	36.075	1.00	33.42
ATOM	1884	N	PRO	A	247	12.525	45.257	31.961	1.00	25.08
ATOM	1885	CA	PRO	A	247	13.885	45.782	31.911	1.00	24.89
ATOM	1886	C	PRO	A	247	13.992	47.144	32.579	1.00	24.95
ATOM	1887	O	PRO	A	247	13.217	48.049	32.297	1.00	24.55
ATOM	1888	CB	PRO	A	247	14.182	45.899	30.413	1.00	25.05
ATOM	1889	CG	PRO	A	247	13.016	45.325	29.702	1.00	25.39
ATOM	1890	CD	PRO	A	247	11.890	45.184	30.640	1.00	25.02
ATOM	1891	N	ASP	A	248	14.976	47.267	33.460	1.00	24.53
ATOM	1892	CA	ASP	A	248	15.216	48.489	34.167	1.00	24.65
ATOM	1893	C	ASP	A	248	16.271	49.284	33.389	1.00	24.54
ATOM	1894	O	ASP	A	248	17.472	49.077	33.552	1.00	23.15
ATOM	1895	CB	ASP	A	248	15.706	48.144	35.564	1.00	24.64
ATOM	1896	CG	ASP	A	248	15.787	49.343	36.454	1.00	26.02
ATOM	1897	OD1	ASP	A	248	16.035	50.466	35.944	1.00	26.95
ATOM	1898	OD2	ASP	A	248	15.609	49.249	37.684	1.00	27.69
ATOM	1899	N	TYR	A	249	15.812	50.181	32.525	1.00	24.81
ATOM	1900	CA	TYR	A	249	16.720	50.936	31.673	1.00	25.45
ATOM	1901	C	TYR	A	249	17.573	51.954	32.426	1.00	25.90
ATOM	1902	O	TYR	A	249	18.585	52.421	31.897	1.00	25.80

N
C
C
O
C
C
C
N
C
C
O
C
C
O
O
N
C
C
C
C
C
C
C
N
C
C
C
O
C
C
O
O
N
C
C
C
C
C
C
N
C
C
C
O
C
C
O
O
N
C
C
C
O

ATOM	1903	CB	TYR	A	249	15.948	51.615	30.544	1.00	25.57
ATOM	1904	CG	TYR	A	249	15.244	50.651	29.630	1.00	24.58
ATOM	1905	CD1	TYR	A	249	15.930	49.624	29.003	1.00	25.44
ATOM	1906	CD2	TYR	A	249	13.890	50.764	29.404	1.00	24.48
ATOM	1907	CE1	TYR	A	249	15.276	48.731	28.173	1.00	25.01
ATOM	1908	CE2	TYR	A	249	13.234	49.892	28.583	1.00	24.43
ATOM	1909	CZ	TYR	A	249	13.927	48.883	27.969	1.00	24.47
ATOM	1910	OH	TYR	A	249	13.252	48.043	27.142	1.00	25.97
ATOM	1911	N	GLU	A	250	17.195	52.278	33.660	1.00	26.10
ATOM	1912	CA	GLU	A	250	17.999	53.190	34.458	1.00	26.93
ATOM	1913	C	GLU	A	250	19.259	52.492	34.929	1.00	25.85
ATOM	1914	O	GLU	A	250	20.329	53.086	34.964	1.00	25.90
ATOM	1915	CB	GLU	A	250	17.219	53.740	35.661	1.00	27.56
ATOM	1916	CG	GLU	A	250	16.021	54.576	35.246	1.00	32.34
ATOM	1917	CD	GLU	A	250	15.420	55.366	36.385	1.00	36.85
ATOM	1918	OE1	GLU	A	250	15.945	55.309	37.516	1.00	42.35
ATOM	1919	OE2	GLU	A	250	14.422	56.057	36.142	1.00	40.82
ATOM	1920	N	ARG	A	251	19.136	51.233	35.315	1.00	24.58
ATOM	1921	CA	ARG	A	251	20.297	50.502	35.756	1.00	23.76
ATOM	1922	C	ARG	A	251	21.030	49.909	34.568	1.00	22.91
ATOM	1923	O	ARG	A	251	22.242	49.773	34.602	1.00	22.42
ATOM	1924	CB	ARG	A	251	19.896	49.374	36.696	1.00	24.55
ATOM	1925	CG	ARG	A	251	19.421	49.802	38.077	1.00	25.88
ATOM	1926	CD	ARG	A	251	18.894	48.629	38.921	1.00	29.50
ATOM	1927	NE	ARG	A	251	19.993	47.764	39.349	1.00	31.63
ATOM	1928	CZ	ARG	A	251	19.894	46.467	39.623	1.00	33.13
ATOM	1929	NH1	ARG	A	251	18.736	45.826	39.521	1.00	33.12
ATOM	1930	NH2	ARG	A	251	20.978	45.805	39.998	1.00	33.93
ATOM	1931	N	PHE	A	252	20.300	49.560	33.515	1.00	21.47
ATOM	1932	CA	PHE	A	252	20.898	48.863	32.379	1.00	21.32
ATOM	1933	C	PHE	A	252	20.576	49.552	31.056	1.00	20.68
ATOM	1934	O	PHE	A	252	19.934	48.984	30.187	1.00	20.79
ATOM	1935	CB	PHE	A	252	20.355	47.434	32.344	1.00	21.09
ATOM	1936	CG	PHE	A	252	20.373	46.738	33.690	1.00	20.96
ATOM	1937	CD1	PHE	A	252	21.555	46.560	34.376	1.00	19.28
ATOM	1938	CD2	PHE	A	252	19.207	46.238	34.247	1.00	21.43
ATOM	1939	CE1	PHE	A	252	21.571	45.891	35.609	1.00	20.54
ATOM	1940	CE2	PHE	A	252	19.217	45.588	35.488	1.00	22.13
ATOM	1941	CZ	PHE	A	252	20.403	45.414	36.156	1.00	20.96
ATOM	1942	N	PRO	A	253	21.057	50.767	30.884	1.00	20.56
ATOM	1943	CA	PRO	A	253	20.658	51.566	29.714	1.00	20.38
ATOM	1944	C	PRO	A	253	20.984	50.916	28.361	1.00	20.20
ATOM	1945	O	PRO	A	253	20.191	51.053	27.428	1.00	19.48
ATOM	1946	CB	PRO	A	253	21.383	52.903	29.928	1.00	19.77
ATOM	1947	CG	PRO	A	253	22.562	52.555	30.895	1.00	20.60
ATOM	1948	CD	PRO	A	253	21.986	51.484	31.782	1.00	20.69
ATOM	1949	N	ASN	A	254	22.091	50.187	28.240	1.00	20.32
ATOM	1950	CA	ASN	A	254	22.409	49.613	26.938	1.00	20.12
ATOM	1951	C	ASN	A	254	21.572	48.393	26.554	1.00	19.96
ATOM	1952	O	ASN	A	254	21.675	47.892	25.441	1.00	19.16
ATOM	1953	CB	ASN	A	254	23.893	49.336	26.784	1.00	20.06
ATOM	1954	CG	ASN	A	254	24.706	50.601	26.697	1.00	21.51
ATOM	1955	OD1	ASN	A	254	25.502	50.896	27.598	1.00	23.96
ATOM	1956	ND2	ASN	A	254	24.514	51.373	25.614	1.00	20.38
ATOM	1957	N	PHE	A	255	20.724	47.922	27.455	1.00	19.81
ATOM	1958	CA	PHE	A	255	19.807	46.858	27.063	1.00	19.54
ATOM	1959	C	PHE	A	255	18.855	47.384	25.977	1.00	19.69
ATOM	1960	O	PHE	A	255	18.194	46.606	25.274	1.00	19.77
ATOM	1961	CB	PHE	A	255	19.023	46.330	28.252	1.00	19.22
ATOM	1962	CG	PHE	A	255	18.269	45.105	27.940	1.00	20.47
ATOM	1963	CD1	PHE	A	255	18.928	43.922	27.736	1.00	20.23

[illegible]

ATOM	1964	CD2	PHE	A	255	16.909	45.148	27.758	1.00	20.13	C
ATOM	1965	CE1	PHE	A	255	18.243	42.786	27.416	1.00	21.65	C
ATOM	1966	CE2	PHE	A	255	16.227	44.023	27.432	1.00	20.73	C
ATOM	1967	CZ	PHE	A	255	16.886	42.842	27.257	1.00	22.63	C
ATOM	1968	N	GLN	A	256	18.790	48.707	25.841	1.00	19.89	N
ATOM	1969	CA	GLN	A	256	17.969	49.345	24.794	1.00	20.24	C
ATOM	1970	C	GLN	A	256	18.595	49.160	23.403	1.00	19.74	C
ATOM	1971	O	GLN	A	256	17.969	49.443	22.384	1.00	19.92	O
ATOM	1972	CB	GLN	A	256	17.778	50.849	25.095	1.00	20.01	C
ATOM	1973	CG	GLN	A	256	16.736	51.095	26.197	1.00	21.20	C
ATOM	1974	CD	GLN	A	256	16.741	52.513	26.764	1.00	22.14	C
ATOM	1975	OE1	GLN	A	256	15.711	53.195	26.741	1.00	23.01	O
ATOM	1976	NE2	GLN	A	256	17.874	52.943	27.298	1.00	21.42	N
ATOM	1977	N	ASN	A	257	19.845	48.716	23.385	1.00	19.66	N
ATOM	1978	CA	ASN	A	257	20.580	48.485	22.155	1.00	20.07	C
ATOM	1979	C	ASN	A	257	20.757	47.005	21.811	1.00	20.73	C
ATOM	1980	O	ASN	A	257	21.416	46.694	20.823	1.00	20.96	O
ATOM	1981	CB	ASN	A	257	21.969	49.100	22.232	1.00	19.17	C
ATOM	1982	CG	ASN	A	257	21.941	50.550	22.635	1.00	20.82	C
ATOM	1983	OD1	ASN	A	257	22.413	50.902	23.722	1.00	21.51	O
ATOM	1984	ND2	ASN	A	257	21.396	51.407	21.763	1.00	16.92	N
ATOM	1985	N	VAL	A	258	20.190	46.098	22.596	1.00	21.59	N
ATOM	1986	CA	VAL	A	258	20.410	44.660	22.347	1.00	23.34	C
ATOM	1987	C	VAL	A	258	19.671	44.172	21.112	1.00	23.85	C
ATOM	1988	O	VAL	A	258	18.549	44.611	20.818	1.00	24.10	O
ATOM	1989	CB	VAL	A	258	20.015	43.806	23.573	1.00	23.68	C
ATOM	1990	CG1	VAL	A	258	18.507	43.720	23.685	1.00	24.12	C
ATOM	1991	CG2	VAL	A	258	20.570	42.436	23.445	1.00	28.30	C
ATOM	1992	N	VAL	A	259	20.314	43.286	20.360	1.00	24.47	N
ATOM	1993	CA	VAL	A	259	19.705	42.721	19.154	1.00	24.91	C
ATOM	1994	C	VAL	A	259	20.004	41.215	19.140	1.00	24.93	C
ATOM	1995	O	VAL	A	259	21.152	40.829	19.119	1.00	25.32	O
ATOM	1996	CB	VAL	A	259	20.281	43.362	17.895	1.00	24.88	C
ATOM	1997	CG1	VAL	A	259	19.667	42.738	16.670	1.00	26.38	C
ATOM	1998	CG2	VAL	A	259	20.051	44.869	17.909	1.00	24.83	C
ATOM	1999	N	GLY	A	260	18.974	40.381	19.186	1.00	25.06	N
ATOM	2000	CA	GLY	A	260	19.166	38.944	19.232	1.00	25.64	C
ATOM	2001	C	GLY	A	260	19.076	38.221	17.887	1.00	25.42	C
ATOM	2002	O	GLY	A	260	18.679	38.796	16.869	1.00	25.77	O
ATOM	2003	N	TYR	A	261	19.497	36.965	17.907	1.00	25.04	N
ATOM	2004	CA	TYR	A	261	19.380	36.049	16.792	1.00	25.30	C
ATOM	2005	C	TYR	A	261	18.428	34.973	17.328	1.00	24.63	C
ATOM	2006	O	TYR	A	261	18.676	34.417	18.392	1.00	24.77	O
ATOM	2007	CB	TYR	A	261	20.729	35.414	16.482	1.00	25.84	C
ATOM	2008	CG	TYR	A	261	21.750	36.315	15.825	1.00	28.57	C
ATOM	2009	CD1	TYR	A	261	21.821	36.423	14.447	1.00	34.41	C
ATOM	2010	CD2	TYR	A	261	22.631	37.049	16.574	1.00	29.87	C
ATOM	2011	CE1	TYR	A	261	22.752	37.255	13.838	1.00	35.64	C
ATOM	2012	CE2	TYR	A	261	23.576	37.852	15.986	1.00	32.26	C
ATOM	2013	CZ	TYR	A	261	23.644	37.949	14.623	1.00	34.92	C
ATOM	2014	OH	TYR	A	261	24.582	38.772	14.047	1.00	38.39	O
ATOM	2015	N	GLU	A	262	17.325	34.698	16.658	1.00	23.64	N
ATOM	2016	CA	GLU	A	262	16.376	33.758	17.244	1.00	24.03	C
ATOM	2017	C	GLU	A	262	15.933	32.660	16.306	1.00	23.15	C
ATOM	2018	O	GLU	A	262	16.063	32.766	15.078	1.00	22.51	O
ATOM	2019	CB	GLU	A	262	15.172	34.482	17.861	1.00	24.11	C
ATOM	2020	CG	GLU	A	262	13.899	34.502	17.057	1.00	26.35	C
ATOM	2021	CD	GLU	A	262	12.744	35.177	17.785	1.00	27.98	C
ATOM	2022	OE1	GLU	A	262	12.743	36.415	17.872	1.00	27.19	O
ATOM	2023	OE2	GLU	A	262	11.818	34.476	18.254	1.00	29.21	O
ATOM	2024	N	THR	A	263	15.455	31.587	16.917	1.00	22.58	N

ATOM	2025	CA	THR	A	263	14.939	30.458	16.170	1.00	22.52
ATOM	2026	C	THR	A	263	14.047	29.617	17.076	1.00	22.48
ATOM	2027	O	THR	A	263	14.107	29.712	18.313	1.00	22.14
ATOM	2028	CB	THR	A	263	16.117	29.607	15.659	1.00	22.55
ATOM	2029	OG1	THR	A	263	15.665	28.666	14.675	1.00	22.25
ATOM	2030	CG2	THR	A	263	16.693	28.743	16.770	1.00	22.79
ATOM	2031	N	VAL	A	264	13.208	28.797	16.472	1.00	21.62
ATOM	2032	CA	VAL	A	264	12.453	27.860	17.263	1.00	22.45
ATOM	2033	C	VAL	A	264	12.871	26.474	16.837	1.00	22.76
ATOM	2034	O	VAL	A	264	12.779	26.134	15.673	1.00	22.77
ATOM	2035	CB	VAL	A	264	10.964	28.027	17.087	1.00	23.08
ATOM	2036	CG1	VAL	A	264	10.234	26.805	17.643	1.00	23.67
ATOM	2037	CG2	VAL	A	264	10.499	29.344	17.767	1.00	22.75
ATOM	2038	N	VAL	A	265	13.389	25.695	17.777	1.00	22.93
ATOM	2039	CA	VAL	A	265	13.779	24.329	17.458	1.00	23.08
ATOM	2040	C	VAL	A	265	12.738	23.317	17.886	1.00	22.86
ATOM	2041	O	VAL	A	265	12.105	23.473	18.932	1.00	23.06
ATOM	2042	CB	VAL	A	265	15.143	23.937	18.075	1.00	22.94
ATOM	2043	CG1	VAL	A	265	16.243	24.745	17.404	1.00	23.66
ATOM	2044	CG2	VAL	A	265	15.169	24.122	19.592	1.00	22.99
ATOM	2045	N	GLY	A	266	12.574	22.290	17.056	1.00	21.77
ATOM	2046	CA	GLY	A	266	11.683	21.178	17.355	1.00	21.98
ATOM	2047	C	GLY	A	266	12.376	19.826	17.263	1.00	20.97
ATOM	2048	O	GLY	A	266	13.562	19.757	16.999	1.00	20.63
ATOM	2049	N	PRO	A	267	11.643	18.737	17.461	1.00	21.23
ATOM	2050	CA	PRO	A	267	12.253	17.403	17.427	1.00	20.68
ATOM	2051	C	PRO	A	267	13.126	17.183	16.203	1.00	20.04
ATOM	2052	O	PRO	A	267	12.666	17.371	15.081	1.00	19.35
ATOM	2053	CB	PRO	A	267	11.039	16.463	17.379	1.00	21.29
ATOM	2054	CG	PRO	A	267	9.970	17.202	18.037	1.00	22.85
ATOM	2055	CD	PRO	A	267	10.188	18.676	17.710	1.00	21.61
ATOM	2056	N	GLY	A	268	14.376	16.787	16.411	1.00	20.31
ATOM	2057	CA	GLY	A	268	15.278	16.518	15.305	1.00	19.95
ATOM	2058	C	GLY	A	268	16.256	17.627	14.993	1.00	20.50
ATOM	2059	O	GLY	A	268	17.277	17.400	14.311	1.00	21.62
ATOM	2060	N	ASP	A	269	15.965	18.831	15.477	1.00	19.81
ATOM	2061	CA	ASP	A	269	16.818	19.977	15.217	1.00	19.75
ATOM	2062	C	ASP	A	269	17.970	20.003	16.215	1.00	19.67
ATOM	2063	O	ASP	A	269	17.811	19.625	17.377	1.00	18.76
ATOM	2064	CB	ASP	A	269	16.032	21.288	15.386	1.00	19.99
ATOM	2065	CG	ASP	A	269	14.916	21.464	14.374	1.00	20.91
ATOM	2066	OD1	ASP	A	269	14.935	20.842	13.287	1.00	22.13
ATOM	2067	OD2	ASP	A	269	13.987	22.259	14.576	1.00	23.27
ATOM	2068	N	VAL	A	270	19.116	20.481	15.746	1.00	19.56
ATOM	2069	CA	VAL	A	270	20.276	20.700	16.595	1.00	19.57
ATOM	2070	C	VAL	A	270	20.747	22.137	16.444	1.00	19.33
ATOM	2071	O	VAL	A	270	21.033	22.594	15.352	1.00	19.78
ATOM	2072	CB	VAL	A	270	21.404	19.776	16.215	1.00	19.55
ATOM	2073	CG1	VAL	A	270	22.646	20.154	16.966	1.00	20.05
ATOM	2074	CG2	VAL	A	270	21.001	18.342	16.535	1.00	19.44
ATOM	2075	N	LEU	A	271	20.786	22.870	17.539	1.00	19.33
ATOM	2076	CA	LEU	A	271	21.263	24.229	17.501	1.00	19.02
ATOM	2077	C	LEU	A	271	22.677	24.272	18.036	1.00	19.15
ATOM	2078	O	LEU	A	271	22.961	23.787	19.130	1.00	18.74
ATOM	2079	CB	LEU	A	271	20.376	25.120	18.331	1.00	19.93
ATOM	2080	CG	LEU	A	271	20.892	26.549	18.573	1.00	20.11
ATOM	2081	CD1	LEU	A	271	21.011	27.358	17.296	1.00	19.41
ATOM	2082	CD2	LEU	A	271	19.979	27.245	19.555	1.00	20.25
ATOM	2083	N	TYR	A	272	23.582	24.818	17.244	1.00	19.15
ATOM	2084	CA	TYR	A	272	24.924	25.051	17.718	1.00	19.70
ATOM	2085	C	TYR	A	272	24.914	26.360	18.483	1.00	19.63

[illegible]

ATOM	2086	O	TYR A 272	24.646	27.395	17.897	1.00	20.44
ATOM	2087	CB	TYR A 272	25.908	25.172	16.547	1.00	19.86
ATOM	2088	CG	TYR A 272	27.293	25.654	16.962	1.00	21.59
ATOM	2089	CD1	TYR A 272	27.952	25.093	18.048	1.00	22.02
ATOM	2090	CD2	TYR A 272	27.936	26.668	16.278	1.00	20.87
ATOM	2091	CE1	TYR A 272	29.214	25.522	18.421	1.00	19.87
ATOM	2092	CE2	TYR A 272	29.200	27.098	16.649	1.00	20.88
ATOM	2093	CZ	TYR A 272	29.834	26.517	17.721	1.00	21.07
ATOM	2094	OH	TYR A 272	31.091	26.937	18.104	1.00	18.93
ATOM	2095	N	ILE A 273	25.210	26.308	19.776	1.00	19.88
ATOM	2096	CA	ILE A 273	25.342	27.500	20.598	1.00	20.12
ATOM	2097	C	ILE A 273	26.837	27.683	20.883	1.00	20.39
ATOM	2098	O	ILE A 273	27.393	27.000	21.760	1.00	19.88
ATOM	2099	CB	ILE A 273	24.576	27.366	21.901	1.00	19.72
ATOM	2100	CG1	ILE A 273	23.111	27.045	21.624	1.00	20.57
ATOM	2101	CG2	ILE A 273	24.661	28.695	22.696	1.00	20.99
ATOM	2102	CD1	ILE A 273	22.296	26.797	22.876	1.00	20.93
ATOM	2103	N	PRO A 274	27.487	28.611	20.182	1.00	20.23
ATOM	2104	CA	PRO A 274	28.938	28.743	20.326	1.00	21.16
ATOM	2105	C	PRO A 274	29.289	29.264	21.704	1.00	21.39
ATOM	2106	O	PRO A 274	28.520	30.027	22.317	1.00	20.89
ATOM	2107	CB	PRO A 274	29.353	29.757	19.230	1.00	21.62
ATOM	2108	CG	PRO A 274	28.089	30.056	18.410	1.00	21.22
ATOM	2109	CD	PRO A 274	26.919	29.621	19.282	1.00	20.51
ATOM	2110	N	MET A 275	30.450	28.831	22.180	1.00	21.53
ATOM	2111	CA	MET A 275	30.953	29.207	23.479	1.00	22.30
ATOM	2112	C	MET A 275	30.920	30.734	23.636	1.00	21.94
ATOM	2113	O	MET A 275	31.160	31.442	22.675	1.00	20.78
ATOM	2114	CB	MET A 275	32.367	28.695	23.589	1.00	23.06
ATOM	2115	CG	MET A 275	32.937	28.734	24.966	1.00	27.00
ATOM	2116	SD	MET A 275	34.545	27.926	24.991	1.00	33.74
ATOM	2117	CE	MET A 275	35.263	28.479	23.499	1.00	32.50
ATOM	2118	N	TYR A 276	30.583	31.211	24.832	1.00	21.96
ATOM	2119	CA	TYR A 276	30.485	32.656	25.138	1.00	23.15
ATOM	2120	C	TYR A 276	29.256	33.365	24.546	1.00	22.39
ATOM	2121	O	TYR A 276	28.989	34.496	24.899	1.00	22.58
ATOM	2122	CB	TYR A 276	31.776	33.409	24.760	1.00	23.80
ATOM	2123	CG	TYR A 276	32.904	33.136	25.729	1.00	28.82
ATOM	2124	CD1	TYR A 276	32.940	33.756	26.951	1.00	32.73
ATOM	2125	CD2	TYR A 276	33.935	32.246	25.415	1.00	35.83
ATOM	2126	CE1	TYR A 276	33.956	33.507	27.864	1.00	35.96
ATOM	2127	CE2	TYR A 276	34.975	31.992	26.336	1.00	38.59
ATOM	2128	CZ	TYR A 276	34.958	32.628	27.562	1.00	38.60
ATOM	2129	OH	TYR A 276	35.951	32.421	28.513	1.00	45.18
ATOM	2130	N	TRP A 277	28.488	32.709	23.684	1.00	21.71
ATOM	2131	CA	TRP A 277	27.281	33.350	23.149	1.00	21.24
ATOM	2132	C	TRP A 277	26.184	33.278	24.152	1.00	20.90
ATOM	2133	O	TRP A 277	25.914	32.217	24.708	1.00	21.96
ATOM	2134	CB	TRP A 277	26.790	32.711	21.856	1.00	21.01
ATOM	2135	CG	TRP A 277	27.547	33.166	20.670	1.00	19.98
ATOM	2136	CD1	TRP A 277	28.875	33.013	20.462	1.00	17.90
ATOM	2137	CD2	TRP A 277	27.030	33.832	19.519	1.00	19.76
ATOM	2138	NE1	TRP A 277	29.218	33.533	19.247	1.00	18.64
ATOM	2139	CE2	TRP A 277	28.106	34.071	18.662	1.00	18.29
ATOM	2140	CE3	TRP A 277	25.764	34.305	19.151	1.00	20.56
ATOM	2141	CZ2	TRP A 277	27.963	34.716	17.442	1.00	19.63
ATOM	2142	CZ3	TRP A 277	25.624	34.950	17.943	1.00	21.16
ATOM	2143	CH2	TRP A 277	26.720	35.137	17.095	1.00	19.92
ATOM	2144	N	TRP A 278	25.574	34.426	24.405	1.00	20.32
ATOM	2145	CA	TRP A 278	24.452	34.533	25.316	1.00	20.13
ATOM	2146	C	TRP A 278	23.268	33.787	24.741	1.00	19.99

ATOM	2147	O	TRP	A	278	23.083	33.758	23.538	1.00	20.42
ATOM	2148	CB	TRP	A	278	24.025	35.999	25.441	1.00	20.65
ATOM	2149	CG	TRP	A	278	25.031	36.864	26.107	1.00	20.56
ATOM	2150	CD1	TRP	A	278	26.078	37.526	25.519	1.00	19.63
ATOM	2151	CD2	TRP	A	278	25.096	37.160	27.501	1.00	19.59
ATOM	2152	NE1	TRP	A	278	26.772	38.233	26.469	1.00	20.70
ATOM	2153	CE2	TRP	A	278	26.191	38.012	27.699	1.00	20.10
ATOM	2154	CE3	TRP	A	278	24.318	36.811	28.601	1.00	20.98
ATOM	2155	CZ2	TRP	A	278	26.527	38.516	28.951	1.00	20.24
ATOM	2156	CZ3	TRP	A	278	24.668	37.299	29.849	1.00	23.26
ATOM	2157	CH2	TRP	A	278	25.759	38.146	30.011	1.00	20.41
ATOM	2158	N	HIS	A	279	22.463	33.194	25.591	1.00	19.84
ATOM	2159	CA	HIS	A	279	21.269	32.530	25.122	1.00	20.78
ATOM	2160	C	HIS	A	279	20.202	32.495	26.192	1.00	20.37
ATOM	2161	O	HIS	A	279	20.479	32.275	27.370	1.00	20.66
ATOM	2162	CB	HIS	A	279	21.540	31.098	24.604	1.00	20.86
ATOM	2163	CG	HIS	A	279	22.349	30.238	25.528	1.00	21.83
ATOM	2164	ND1	HIS	A	279	23.720	30.324	25.608	1.00	23.19
ATOM	2165	CD2	HIS	A	279	21.988	29.240	26.371	1.00	23.96
ATOM	2166	CE1	HIS	A	279	24.175	29.411	26.449	1.00	22.20
ATOM	2167	NE2	HIS	A	279	23.144	28.747	26.940	1.00	23.84
ATOM	2168	N	HIS	A	280	18.994	32.718	25.711	1.00	20.90
ATOM	2169	CA	HIS	A	280	17.742	32.758	26.441	1.00	21.22
ATOM	2170	C	HIS	A	280	16.910	31.676	25.793	1.00	21.42
ATOM	2171	O	HIS	A	280	16.749	31.685	24.580	1.00	21.70
ATOM	2172	CB	HIS	A	280	17.099	34.124	26.224	1.00	21.04
ATOM	2173	CG	HIS	A	280	15.632	34.168	26.489	1.00	21.54
ATOM	2174	ND1	HIS	A	280	15.096	34.836	27.571	1.00	22.76
ATOM	2175	CD2	HIS	A	280	14.581	33.660	25.802	1.00	22.74
ATOM	2176	CE1	HIS	A	280	13.779	34.721	27.546	1.00	23.23
ATOM	2177	NE2	HIS	A	280	13.441	34.014	26.482	1.00	23.05
ATOM	2178	N	ILE	A	281	16.367	30.755	26.584	1.00	22.31
ATOM	2179	CA	ILE	A	281	15.649	29.599	26.040	1.00	22.39
ATOM	2180	C	ILE	A	281	14.304	29.443	26.722	1.00	22.72
ATOM	2181	O	ILE	A	281	14.216	29.419	27.945	1.00	22.70
ATOM	2182	CB	ILE	A	281	16.527	28.336	26.189	1.00	22.98
ATOM	2183	CG1	ILE	A	281	17.771	28.504	25.305	1.00	24.90
ATOM	2184	CG2	ILE	A	281	15.770	27.059	25.771	1.00	21.60
ATOM	2185	CD1	ILE	A	281	18.795	27.556	25.554	1.00	27.31
ATOM	2186	N	GLU	A	282	13.252	29.350	25.920	1.00	22.93
ATOM	2187	CA	GLU	A	282	11.903	29.205	26.458	1.00	23.35
ATOM	2188	C	GLU	A	282	11.101	28.105	25.769	1.00	22.59
ATOM	2189	O	GLU	A	282	11.092	27.982	24.549	1.00	21.77
ATOM	2190	CB	GLU	A	282	11.144	30.538	26.393	1.00	23.47
ATOM	2191	CG	GLU	A	282	10.830	31.055	25.009	1.00	25.70
ATOM	2192	CD	GLU	A	282	10.281	32.483	25.023	1.00	25.79
ATOM	2193	OE1	GLU	A	282	10.898	33.356	25.665	1.00	27.49
ATOM	2194	OE2	GLU	A	282	9.241	32.740	24.391	1.00	26.03
ATOM	2195	N	SER	A	283	10.456	27.293	26.588	1.00	22.44
ATOM	2196	CA	SER	A	283	9.570	26.246	26.103	1.00	23.00
ATOM	2197	C	SER	A	283	8.256	26.916	25.753	1.00	23.56
ATOM	2198	O	SER	A	283	7.685	27.615	26.584	1.00	24.14
ATOM	2199	CB	SER	A	283	9.346	25.204	27.188	1.00	22.59
ATOM	2200	OG	SER	A	283	10.496	24.400	27.345	1.00	22.67
ATOM	2201	N	LEU	A	284	7.763	26.694	24.541	1.00	24.07
ATOM	2202	CA	LEU	A	284	6.581	27.412	24.070	1.00	24.80
ATOM	2203	C	LEU	A	284	5.386	27.324	25.004	1.00	24.51
ATOM	2204	O	LEU	A	284	5.098	26.271	25.602	1.00	23.93
ATOM	2205	CB	LEU	A	284	6.161	26.947	22.682	1.00	25.09
ATOM	2206	CG	LEU	A	284	7.185	27.083	21.555	1.00	26.46
ATOM	2207	CD1	LEU	A	284	6.475	27.236	20.215	1.00	28.29

O
C
C
C
C
N
C
C
C
C
C
N
C
C
C
C
N
N
N
C
C
O
C
C
C
N
C
C
C
C
C
O
C
C
O
N
C
C
C
O
N
C
C
C
C
C

ATOM	2208	CD2	LEU	A	284	8.123	28.188	21.775	1.00	26.10
ATOM	2209	N	LEU	A	285	4.718	28.463	25.136	1.00	24.41
ATOM	2210	CA	LEU	A	285	3.506	28.554	25.930	1.00	24.83
ATOM	2211	C	LEU	A	285	2.526	27.583	25.323	1.00	24.68
ATOM	2212	O	LEU	A	285	2.393	27.516	24.110	1.00	24.08
ATOM	2213	CB	LEU	A	285	2.927	29.961	25.872	1.00	24.79
ATOM	2214	CG	LEU	A	285	3.825	31.078	26.397	1.00	25.19
ATOM	2215	CD1	LEU	A	285	3.298	32.434	25.954	1.00	26.02
ATOM	2216	CD2	LEU	A	285	3.925	30.999	27.898	1.00	26.27
ATOM	2217	N	ASN	A	286	1.867	26.815	26.173	1.00	25.20
ATOM	2218	CA	ASN	A	286	0.867	25.848	25.730	1.00	25.85
ATOM	2219	C	ASN	A	286	1.371	24.791	24.742	1.00	25.23
ATOM	2220	O	ASN	A	286	0.594	24.267	23.965	1.00	24.38
ATOM	2221	CB	ASN	A	286	-0.319	26.599	25.124	1.00	26.54
ATOM	2222	CG	ASN	A	286	-0.957	27.546	26.109	1.00	29.40
ATOM	2223	OD1	ASN	A	286	-1.478	27.118	27.140	1.00	33.49
ATOM	2224	ND2	ASN	A	286	-0.893	28.843	25.821	1.00	33.58
ATOM	2225	N	GLY	A	287	2.669	24.487	24.774	1.00	24.55
ATOM	2226	CA	GLY	A	287	3.243	23.518	23.863	1.00	24.10
ATOM	2227	C	GLY	A	287	3.524	22.167	24.487	1.00	23.80
ATOM	2228	O	GLY	A	287	4.110	21.301	23.838	1.00	24.10
ATOM	2229	N	GLY	A	288	3.100	21.971	25.734	1.00	23.32
ATOM	2230	CA	GLY	A	288	3.360	20.730	26.439	1.00	23.24
ATOM	2231	C	GLY	A	288	4.808	20.673	26.903	1.00	23.49
ATOM	2232	O	GLY	A	288	5.548	21.646	26.741	1.00	22.86
ATOM	2233	N	ILE	A	289	5.220	19.539	27.462	1.00	23.36
ATOM	2234	CA	ILE	A	289	6.571	19.406	27.983	1.00	23.99
ATOM	2235	C	ILE	A	289	7.601	19.374	26.874	1.00	23.28
ATOM	2236	O	ILE	A	289	7.324	18.958	25.755	1.00	23.68
ATOM	2237	CB	ILE	A	289	6.750	18.123	28.817	1.00	24.68
ATOM	2238	CG1	ILE	A	289	6.819	16.912	27.901	1.00	26.90
ATOM	2239	CG2	ILE	A	289	5.648	17.983	29.885	1.00	25.75
ATOM	2240	CD1	ILE	A	289	7.294	15.640	28.613	1.00	29.37
ATOM	2241	N	THR	A	290	8.811	19.790	27.210	1.00	22.42
ATOM	2242	CA	THR	A	290	9.904	19.768	26.264	1.00	21.38
ATOM	2243	C	THR	A	290	11.030	18.880	26.774	1.00	20.56
ATOM	2244	O	THR	A	290	11.243	18.743	27.986	1.00	19.81
ATOM	2245	CB	THR	A	290	10.461	21.175	26.053	1.00	21.70
ATOM	2246	OG1	THR	A	290	10.823	21.761	27.308	1.00	20.15
ATOM	2247	CG2	THR	A	290	9.405	22.108	25.460	1.00	21.76
ATOM	2248	N	ILE	A	291	11.757	18.296	25.841	1.00	19.36
ATOM	2249	CA	ILE	A	291	12.903	17.503	26.190	1.00	19.73
ATOM	2250	C	ILE	A	291	14.044	17.865	25.286	1.00	19.29
ATOM	2251	O	ILE	A	291	13.862	17.976	24.083	1.00	19.81
ATOM	2252	CB	ILE	A	291	12.598	15.992	26.085	1.00	20.08
ATOM	2253	CG1	ILE	A	291	11.467	15.609	27.044	1.00	20.06
ATOM	2254	CG2	ILE	A	291	13.873	15.214	26.378	1.00	20.21
ATOM	2255	CD1	ILE	A	291	11.028	14.131	26.931	1.00	21.77
ATOM	2256	N	THR	A	292	15.215	18.070	25.882	1.00	18.95
ATOM	2257	CA	THR	A	292	16.406	18.407	25.156	1.00	19.17
ATOM	2258	C	THR	A	292	17.589	17.691	25.753	1.00	19.18
ATOM	2259	O	THR	A	292	17.671	17.497	26.965	1.00	19.21
ATOM	2260	CB	THR	A	292	16.703	19.937	25.259	1.00	19.83
ATOM	2261	OG1	THR	A	292	15.559	20.724	24.879	1.00	20.23
ATOM	2262	CG2	THR	A	292	17.787	20.350	24.270	1.00	20.15
ATOM	2263	N	VAL	A	293	18.530	17.314	24.903	1.00	19.25
ATOM	2264	CA	VAL	A	293	19.809	16.824	25.387	1.00	19.51
ATOM	2265	C	VAL	A	293	20.912	17.667	24.749	1.00	19.58
ATOM	2266	O	VAL	A	293	20.962	17.792	23.535	1.00	18.47
ATOM	2267	CB	VAL	A	293	20.024	15.345	25.063	1.00	19.73
ATOM	2268	CG1	VAL	A	293	21.490	14.951	25.250	1.00	20.24

ATOM	2269	CG2	VAL	A	293	19.146	14.478	25.968	1.00	20.65
ATOM	2270	N	ASN	A	294	21.766	18.277	25.569	1.00	19.85
ATOM	2271	CA	ASN	A	294	22.882	19.057	25.039	1.00	20.26
ATOM	2272	C	ASN	A	294	24.179	18.232	25.052	1.00	20.35
ATOM	2273	O	ASN	A	294	24.233	17.131	25.634	1.00	21.82
ATOM	2274	CB	ASN	A	294	23.008	20.443	25.727	1.00	20.82
ATOM	2275	CG	ASN	A	294	23.676	20.388	27.102	1.00	21.84
ATOM	2276	OD1	ASN	A	294	24.209	19.354	27.495	1.00	21.66
ATOM	2277	ND2	ASN	A	294	23.630	21.519	27.849	1.00	21.23
ATOM	2278	N	PHE	A	295	25.186	18.729	24.346	1.00	20.69
ATOM	2279	CA	PHE	A	295	26.490	18.095	24.233	1.00	20.46
ATOM	2280	C	PHE	A	295	27.452	19.240	24.494	1.00	21.42
ATOM	2281	O	PHE	A	295	27.573	20.167	23.653	1.00	20.85
ATOM	2282	CB	PHE	A	295	26.728	17.569	22.814	1.00	20.62
ATOM	2283	CG	PHE	A	295	25.898	16.366	22.440	1.00	21.15
ATOM	2284	CD1	PHE	A	295	24.526	16.468	22.247	1.00	21.98
ATOM	2285	CD2	PHE	A	295	26.498	15.130	22.256	1.00	19.31
ATOM	2286	CE1	PHE	A	295	23.791	15.343	21.902	1.00	19.77
ATOM	2287	CE2	PHE	A	295	25.762	14.037	21.885	1.00	17.96
ATOM	2288	CZ	PHE	A	295	24.424	14.138	21.720	1.00	17.87
ATOM	2289	N	TRP	A	296	28.123	19.196	25.645	1.00	21.19
ATOM	2290	CA	TRP	A	296	29.023	20.275	26.050	1.00	21.46
ATOM	2291	C	TRP	A	296	30.481	19.834	25.873	1.00	21.42
ATOM	2292	O	TRP	A	296	30.898	18.795	26.387	1.00	21.96
ATOM	2293	CB	TRP	A	296	28.760	20.669	27.498	1.00	21.13
ATOM	2294	CG	TRP	A	296	27.853	21.859	27.710	1.00	21.95
ATOM	2295	CD1	TRP	A	296	27.797	22.987	26.955	1.00	23.23
ATOM	2296	CD2	TRP	A	296	26.935	22.067	28.797	1.00	22.27
ATOM	2297	NE1	TRP	A	296	26.882	23.869	27.478	1.00	22.26
ATOM	2298	CE2	TRP	A	296	26.336	23.329	28.607	1.00	22.53
ATOM	2299	CE3	TRP	A	296	26.531	21.298	29.894	1.00	24.83
ATOM	2300	CZ2	TRP	A	296	25.362	23.840	29.463	1.00	22.34
ATOM	2301	CZ3	TRP	A	296	25.557	21.810	30.754	1.00	24.59
ATOM	2302	CH2	TRP	A	296	24.993	23.075	30.531	1.00	23.75
ATOM	2303	N	TYR	A	297	31.235	20.632	25.126	1.00	21.64
ATOM	2304	CA	TYR	A	297	32.633	20.361	24.845	1.00	21.81
ATOM	2305	C	TYR	A	297	33.482	21.523	25.320	1.00	22.41
ATOM	2306	O	TYR	A	297	33.110	22.682	25.146	1.00	21.78
ATOM	2307	CB	TYR	A	297	32.862	20.198	23.344	1.00	21.49
ATOM	2308	CG	TYR	A	297	32.287	18.943	22.761	1.00	21.77
ATOM	2309	CD1	TYR	A	297	30.934	18.870	22.415	1.00	21.68
ATOM	2310	CD2	TYR	A	297	33.083	17.841	22.526	1.00	19.80
ATOM	2311	CE1	TYR	A	297	30.405	17.734	21.890	1.00	22.16
ATOM	2312	CE2	TYR	A	297	32.555	16.690	21.993	1.00	20.99
ATOM	2313	CZ	TYR	A	297	31.215	16.637	21.686	1.00	21.46
ATOM	2314	OH	TYR	A	297	30.673	15.494	21.159	1.00	22.11
ATOM	2315	N	LYS	A	298	34.632	21.220	25.899	1.00	23.17
ATOM	2316	CA	LYS	A	298	35.566	22.273	26.288	1.00	24.45
ATOM	2317	C	LYS	A	298	36.074	22.912	25.021	1.00	24.21
ATOM	2318	O	LYS	A	298	36.264	22.228	24.025	1.00	23.35
ATOM	2319	CB	LYS	A	298	36.748	21.699	27.085	1.00	24.85
ATOM	2320	CG	LYS	A	298	36.436	21.416	28.552	1.00	28.00
ATOM	2321	CD	LYS	A	298	37.657	20.865	29.298	1.00	31.59
ATOM	2322	CE	LYS	A	298	37.309	20.447	30.730	1.00	33.77
ATOM	2323	NZ	LYS	A	298	38.406	19.627	31.367	1.00	36.02
ATOM	2324	N	GLY	A	299	36.317	24.216	25.056	1.00	25.28
ATOM	2325	CA	GLY	A	299	36.818	24.909	23.886	1.00	26.38
ATOM	2326	C	GLY	A	299	38.254	24.542	23.547	1.00	27.78
ATOM	2327	O	GLY	A	299	38.935	23.895	24.315	1.00	27.16
ATOM	2328	N	ALA	A	300	38.690	24.949	22.366	1.00	30.42
ATOM	2329	CA	ALA	A	300	40.062	24.762	21.918	1.00	32.93

[illegible]

ATOM	2330	C	ALA	A	300	41.021	25.544	22.819	1.00	35.12
ATOM	2331	O	ALA	A	300	40.597	26.337	23.646	1.00	34.87
ATOM	2332	CB	ALA	A	300	40.199	25.249	20.472	1.00	32.98
ATOM	2333	N	PRO	A	301	42.318	25.355	22.615	1.00	38.54
ATOM	2334	CA	PRO	A	301	43.347	26.012	23.440	1.00	40.37
ATOM	2335	C	PRO	A	301	43.548	27.484	23.102	1.00	41.99
ATOM	2336	O	PRO	A	301	43.379	27.853	21.950	1.00	42.88
ATOM	2337	CB	PRO	A	301	44.615	25.246	23.061	1.00	40.42
ATOM	2338	CG	PRO	A	301	44.397	24.879	21.638	1.00	39.87
ATOM	2339	CD	PRO	A	301	42.919	24.544	21.535	1.00	38.98
ATOM	2340	N	THR	A	302	43.915	28.293	24.090	1.00	44.60
ATOM	2341	CA	THR	A	302	44.209	29.725	23.912	1.00	46.30
ATOM	2342	C	THR	A	302	45.593	29.790	23.315	1.00	47.41
ATOM	2343	O	THR	A	302	46.534	29.388	23.992	1.00	47.96
ATOM	2344	CB	THR	A	302	44.242	30.391	25.294	1.00	46.57
ATOM	2345	OG1	THR	A	302	42.941	30.320	25.895	1.00	48.00
ATOM	2346	CG2	THR	A	302	44.526	31.869	25.199	1.00	47.24
ATOM	2347	N	PRO	A	303	45.782	30.336	22.112	1.00	48.55
ATOM	2348	CA	PRO	A	303	47.090	30.170	21.473	1.00	48.80
ATOM	2349	C	PRO	A	303	48.210	30.717	22.341	1.00	48.76
ATOM	2350	O	PRO	A	303	47.874	31.450	23.269	1.00	49.07
ATOM	2351	CB	PRO	A	303	46.967	30.980	20.185	1.00	49.07
ATOM	2352	CG	PRO	A	303	45.504	31.101	19.952	1.00	48.93
ATOM	2353	CD	PRO	A	303	44.916	31.243	21.330	1.00	48.62
ATOM	2354	N	GLU	A	307	46.795	36.776	18.436	1.00	52.62
ATOM	2355	CA	GLU	A	307	46.885	37.814	17.415	1.00	52.76
ATOM	2356	C	GLU	A	307	45.865	38.906	17.636	1.00	52.00
ATOM	2357	O	GLU	A	307	44.757	38.658	18.096	1.00	52.48
ATOM	2358	CB	GLU	A	307	46.686	37.246	15.996	1.00	53.32
ATOM	2359	CG	GLU	A	307	46.893	38.307	14.908	1.00	54.63
ATOM	2360	CD	GLU	A	307	46.862	37.764	13.487	1.00	56.43
ATOM	2361	OE1	GLU	A	307	46.527	36.574	13.290	1.00	57.49
ATOM	2362	OE2	GLU	A	307	47.173	38.543	12.558	1.00	57.89
ATOM	2363	N	TYR	A	308	46.255	40.125	17.303	1.00	51.19
ATOM	2364	CA	TYR	A	308	45.367	41.267	17.405	1.00	50.55
ATOM	2365	C	TYR	A	308	44.747	41.494	16.039	1.00	49.51
ATOM	2366	O	TYR	A	308	45.300	41.053	15.028	1.00	49.70
ATOM	2367	CB	TYR	A	308	46.151	42.485	17.867	1.00	50.82
ATOM	2368	CG	TYR	A	308	46.702	42.286	19.259	1.00	52.47
ATOM	2369	CD1	TYR	A	308	45.949	42.632	20.373	1.00	53.15
ATOM	2370	CD2	TYR	A	308	47.948	41.696	19.462	1.00	53.79
ATOM	2371	CE1	TYR	A	308	46.426	42.426	21.643	1.00	54.22
ATOM	2372	CE2	TYR	A	308	48.437	41.487	20.736	1.00	54.76
ATOM	2373	CZ	TYR	A	308	47.670	41.857	21.824	1.00	55.10
ATOM	2374	OH	TYR	A	308	48.146	41.659	23.101	1.00	56.68
ATOM	2375	N	PRO	A	309	43.584	42.135	15.987	1.00	47.88
ATOM	2376	CA	PRO	A	309	42.843	42.621	17.169	1.00	46.21
ATOM	2377	C	PRO	A	309	42.139	41.503	17.952	1.00	43.75
ATOM	2378	O	PRO	A	309	41.768	40.503	17.369	1.00	44.10
ATOM	2379	CB	PRO	A	309	41.804	43.566	16.562	1.00	46.39
ATOM	2380	CG	PRO	A	309	41.610	43.069	15.128	1.00	47.65
ATOM	2381	CD	PRO	A	309	42.899	42.431	14.716	1.00	48.06
ATOM	2382	N	LEU	A	310	41.973	41.672	19.256	1.00	40.94
ATOM	2383	CA	LEU	A	310	41.349	40.650	20.093	1.00	38.75
ATOM	2384	C	LEU	A	310	39.863	40.491	19.817	1.00	36.87
ATOM	2385	O	LEU	A	310	39.148	41.474	19.610	1.00	37.02
ATOM	2386	CB	LEU	A	310	41.499	41.017	21.566	1.00	38.63
ATOM	2387	CG	LEU	A	310	42.571	40.351	22.435	1.00	37.97
ATOM	2388	CD1	LEU	A	310	43.840	40.049	21.711	1.00	37.27
ATOM	2389	CD2	LEU	A	310	42.836	41.234	23.637	1.00	37.27
ATOM	2390	N	LYS	A	311	39.392	39.254	19.847	1.00	34.04

ATOM	2391	CA	LYS	A	311	37.972	38.998	19.702	1.00	32.20
ATOM	2392	C	LYS	A	311	37.208	39.376	20.968	1.00	30.33
ATOM	2393	O	LYS	A	311	37.760	39.421	22.072	1.00	29.17
ATOM	2394	CB	LYS	A	311	37.733	37.564	19.371	1.00	32.59
ATOM	2395	N	ALA	A	312	35.924	39.641	20.788	1.00	28.23
ATOM	2396	CA	ALA	A	312	35.074	40.011	21.895	1.00	27.38
ATOM	2397	C	ALA	A	312	35.148	38.973	23.005	1.00	26.28
ATOM	2398	O	ALA	A	312	35.271	39.334	24.172	1.00	24.54
ATOM	2399	CB	ALA	A	312	33.641	40.205	21.421	1.00	27.09
ATOM	2400	N	HIS	A	313	35.101	37.689	22.649	1.00	25.72
ATOM	2401	CA	HIS	A	313	35.086	36.669	23.674	1.00	25.82
ATOM	2402	C	HIS	A	313	36.399	36.609	24.428	1.00	24.79
ATOM	2403	O	HIS	A	313	36.428	36.183	25.557	1.00	24.19
ATOM	2404	CB	HIS	A	313	34.688	35.293	23.129	1.00	26.92
ATOM	2405	CG	HIS	A	313	35.741	34.641	22.303	1.00	29.84
ATOM	2406	ND1	HIS	A	313	35.896	34.903	20.954	1.00	35.52
ATOM	2407	CD2	HIS	A	313	36.702	33.746	22.627	1.00	33.54
ATOM	2408	CE1	HIS	A	313	36.921	34.208	20.489	1.00	35.03
ATOM	2409	NE2	HIS	A	313	37.424	33.491	21.481	1.00	35.78
ATOM	2410	N	GLN	A	314	37.479	37.042	23.803	1.00	24.43
ATOM	2411	CA	GLN	A	314	38.762	37.092	24.465	1.00	24.34
ATOM	2412	C	GLN	A	314	38.762	38.226	25.510	1.00	24.39
ATOM	2413	O	GLN	A	314	39.327	38.081	26.590	1.00	24.82
ATOM	2414	CB	GLN	A	314	39.882	37.290	23.439	1.00	24.47
ATOM	2415	CG	GLN	A	314	40.032	36.106	22.472	1.00	25.49
ATOM	2416	CD	GLN	A	314	41.036	36.362	21.366	1.00	25.52
ATOM	2417	OE1	GLN	A	314	40.878	37.287	20.563	1.00	27.59
ATOM	2418	NE2	GLN	A	314	42.078	35.553	21.330	1.00	28.22
ATOM	2419	N	LYS	A	315	38.113	39.337	25.196	1.00	23.51
ATOM	2420	CA	LYS	A	315	38.000	40.423	26.154	1.00	23.79
ATOM	2421	C	LYS	A	315	37.125	39.979	27.325	1.00	23.05
ATOM	2422	O	LYS	A	315	37.373	40.347	28.465	1.00	21.38
ATOM	2423	CB	LYS	A	315	37.421	41.667	25.504	1.00	24.09
ATOM	2424	CG	LYS	A	315	38.382	42.286	24.533	1.00	26.96
ATOM	2425	CD	LYS	A	315	37.849	43.552	23.947	1.00	31.28
ATOM	2426	CE	LYS	A	315	38.856	44.156	22.977	1.00	34.34
ATOM	2427	NZ	LYS	A	315	38.207	45.098	22.005	1.00	36.25
ATOM	2428	N	VAL	A	316	36.109	39.177	27.041	1.00	22.30
ATOM	2429	CA	VAL	A	316	35.276	38.673	28.113	1.00	22.39
ATOM	2430	C	VAL	A	316	36.124	37.800	29.063	1.00	22.29
ATOM	2431	O	VAL	A	316	36.040	37.932	30.274	1.00	21.27
ATOM	2432	CB	VAL	A	316	34.065	37.887	27.595	1.00	22.03
ATOM	2433	CG1	VAL	A	316	33.309	37.282	28.750	1.00	21.64
ATOM	2434	CG2	VAL	A	316	33.123	38.796	26.802	1.00	22.46
ATOM	2435	N	ALA	A	317	36.964	36.941	28.499	1.00	22.09
ATOM	2436	CA	ALA	A	317	37.848	36.086	29.295	1.00	21.98
ATOM	2437	C	ALA	A	317	38.783	36.916	30.164	1.00	21.61
ATOM	2438	O	ALA	A	317	39.042	36.573	31.300	1.00	21.05
ATOM	2439	CB	ALA	A	317	38.668	35.147	28.380	1.00	22.01
ATOM	2440	N	ILE	A	318	39.273	38.016	29.606	1.00	21.45
ATOM	2441	CA	ILE	A	318	40.162	38.917	30.318	1.00	21.24
ATOM	2442	C	ILE	A	318	39.431	39.511	31.524	1.00	21.38
ATOM	2443	O	ILE	A	318	39.937	39.462	32.646	1.00	20.76
ATOM	2444	CB	ILE	A	318	40.709	40.023	29.377	1.00	20.58
ATOM	2445	CG1	ILE	A	318	41.715	39.434	28.400	1.00	21.19
ATOM	2446	CG2	ILE	A	318	41.354	41.167	30.185	1.00	20.20
ATOM	2447	CD1	ILE	A	318	42.223	40.419	27.353	1.00	21.35
ATOM	2448	N	MET	A	319	38.234	40.030	31.287	1.00	20.83
ATOM	2449	CA	MET	A	319	37.446	40.622	32.355	1.00	21.31
ATOM	2450	C	MET	A	319	37.154	39.617	33.454	1.00	21.52
ATOM	2451	O	MET	A	319	37.326	39.911	34.651	1.00	21.13

[illegible]

ATOM	2452	CB	MET	A	319	36.177	41.260	31.818	1.00	21.01
ATOM	2453	CG	MET	A	319	36.423	42.519	30.971	1.00	21.17
ATOM	2454	SD	MET	A	319	34.860	43.346	30.459	1.00	22.71
ATOM	2455	CE	MET	A	319	34.204	42.152	29.206	1.00	22.13
ATOM	2456	N	ARG	A	320	36.767	38.408	33.068	1.00	21.34
ATOM	2457	CA	ARG	A	320	36.532	37.399	34.075	1.00	21.12
ATOM	2458	C	ARG	A	320	37.801	37.186	34.898	1.00	21.25
ATOM	2459	O	ARG	A	320	37.747	37.064	36.122	1.00	21.48
ATOM	2460	CB	ARG	A	320	36.102	36.095	33.427	1.00	21.01
ATOM	2461	CG	ARG	A	320	34.723	36.145	32.741	1.00	20.63
ATOM	2462	CD	ARG	A	320	34.324	34.795	32.142	1.00	18.73
ATOM	2463	NE	ARG	A	320	34.225	33.824	33.225	1.00	19.25
ATOM	2464	CZ	ARG	A	320	33.247	33.809	34.115	1.00	19.86
ATOM	2465	NH1	ARG	A	320	32.259	34.689	34.062	1.00	21.08
ATOM	2466	NH2	ARG	A	320	33.263	32.923	35.081	1.00	23.29
ATOM	2467	N	ASN	A	321	38.947	37.123	34.231	1.00	20.94
ATOM	2468	CA	ASN	A	321	40.184	36.855	34.939	1.00	20.88
ATOM	2469	C	ASN	A	321	40.535	37.981	35.916	1.00	20.93
ATOM	2470	O	ASN	A	321	40.962	37.716	37.047	1.00	21.77
ATOM	2471	CB	ASN	A	321	41.324	36.552	33.958	1.00	21.04
ATOM	2472	CG	ASN	A	321	41.288	35.099	33.447	1.00	22.80
ATOM	2473	OD1	ASN	A	321	40.924	34.193	34.185	1.00	23.97
ATOM	2474	ND2	ASN	A	321	41.666	34.887	32.193	1.00	21.35
ATOM	2475	N	ILE	A	322	40.355	39.233	35.505	1.00	20.14
ATOM	2476	CA	ILE	A	322	40.633	40.336	36.408	1.00	20.06
ATOM	2477	C	ILE	A	322	39.742	40.196	37.650	1.00	19.42
ATOM	2478	O	ILE	A	322	40.207	40.345	38.767	1.00	18.73
ATOM	2479	CB	ILE	A	322	40.372	41.690	35.715	1.00	20.21
ATOM	2480	CG1	ILE	A	322	41.320	41.894	34.535	1.00	21.54
ATOM	2481	CG2	ILE	A	322	40.504	42.823	36.699	1.00	20.53
ATOM	2482	CD1	ILE	A	322	42.806	41.798	34.868	1.00	24.56
ATOM	2483	N	GLU	A	323	38.458	39.904	37.454	1.00	18.94
ATOM	2484	CA	GLU	A	323	37.553	39.757	38.576	1.00	18.98
ATOM	2485	C	GLU	A	323	38.027	38.651	39.492	1.00	19.33
ATOM	2486	O	GLU	A	323	38.084	38.832	40.707	1.00	19.05
ATOM	2487	CB	GLU	A	323	36.113	39.538	38.106	1.00	19.25
ATOM	2488	CG	GLU	A	323	35.518	40.803	37.484	1.00	19.34
ATOM	2489	CD	GLU	A	323	34.143	40.616	36.855	1.00	18.71
ATOM	2490	OE1	GLU	A	323	33.183	40.332	37.573	1.00	19.19
ATOM	2491	OE2	GLU	A	323	34.024	40.805	35.636	1.00	19.56
ATOM	2492	N	LYS	A	324	38.425	37.528	38.908	1.00	19.55
ATOM	2493	CA	LYS	A	324	38.874	36.379	39.693	1.00	20.24
ATOM	2494	C	LYS	A	324	40.115	36.707	40.513	1.00	20.68
ATOM	2495	O	LYS	A	324	40.157	36.405	41.699	1.00	20.83
ATOM	2496	CB	LYS	A	324	39.175	35.175	38.789	1.00	20.07
ATOM	2497	CG	LYS	A	324	37.924	34.507	38.212	1.00	20.57
ATOM	2498	CD	LYS	A	324	38.331	33.318	37.350	1.00	20.28
ATOM	2499	CE	LYS	A	324	37.137	32.689	36.621	1.00	19.29
ATOM	2500	NZ	LYS	A	324	37.596	31.460	35.876	1.00	17.17
ATOM	2501	N	MET	A	325	41.112	37.320	39.885	1.00	21.21
ATOM	2502	CA	MET	A	325	42.366	37.657	40.568	1.00	22.87
ATOM	2503	C	MET	A	325	42.162	38.665	41.699	1.00	22.44
ATOM	2504	O	MET	A	325	42.763	38.560	42.771	1.00	22.08
ATOM	2505	CB	MET	A	325	43.380	38.232	39.572	1.00	23.32
ATOM	2506	CG	MET	A	325	43.993	37.195	38.671	1.00	27.47
ATOM	2507	SD	MET	A	325	44.795	37.924	37.197	1.00	35.56
ATOM	2508	CE	MET	A	325	45.664	39.189	38.009	1.00	34.71
ATOM	2509	N	LEU	A	326	41.309	39.641	41.452	1.00	22.71
ATOM	2510	CA	LEU	A	326	41.043	40.670	42.449	1.00	23.24
ATOM	2511	C	LEU	A	326	40.385	40.073	43.679	1.00	23.15
ATOM	2512	O	LEU	A	326	40.735	40.417	44.795	1.00	22.40

C
C
S
C
N
C
C
O
C
C
N
C
N
N
N
C
C
O
C
O
N
N
C
C
C
C
C
O
C
C
C
O
N
C
C
O
C
C
C
C
N
N
C
C
C
O
C
C
S
C
N
C
C
C
O

ATOM	2513	CB	LEU A 326	40.151	41.735	41.865	1.00	23.01
ATOM	2514	CG	LEU A 326	40.677	43.139	41.625	1.00	25.86
ATOM	2515	CD1	LEU A 326	42.187	43.277	41.529	1.00	26.53
ATOM	2516	CD2	LEU A 326	39.986	43.672	40.377	1.00	25.54
ATOM	2517	N	GLY A 327	39.441	39.164	43.460	1.00	23.29
ATOM	2518	CA	GLY A 327	38.760	38.489	44.547	1.00	23.97
ATOM	2519	C	GLY A 327	39.728	37.723	45.418	1.00	24.63
ATOM	2520	O	GLY A 327	39.659	37.783	46.649	1.00	24.75
ATOM	2521	N	GLU A 328	40.644	37.002	44.778	1.00	25.03
ATOM	2522	CA	GLU A 328	41.671	36.265	45.506	1.00	26.08
ATOM	2523	C	GLU A 328	42.661	37.180	46.223	1.00	25.43
ATOM	2524	O	GLU A 328	43.029	36.926	47.367	1.00	25.16
ATOM	2525	CB	GLU A 328	42.442	35.368	44.546	1.00	26.76
ATOM	2526	CG	GLU A 328	41.576	34.278	43.947	1.00	30.36
ATOM	2527	CD	GLU A 328	41.719	32.957	44.676	1.00	35.76
ATOM	2528	OE1	GLU A 328	42.091	32.986	45.878	1.00	38.79
ATOM	2529	OE2	GLU A 328	41.483	31.896	44.034	1.00	38.82
ATOM	2530	N	ALA A 329	43.094	38.240	45.552	1.00	24.91
ATOM	2531	CA	ALA A 329	44.102	39.119	46.134	1.00	24.95
ATOM	2532	C	ALA A 329	43.535	39.929	47.285	1.00	24.94
ATOM	2533	O	ALA A 329	44.197	40.147	48.276	1.00	24.57
ATOM	2534	CB	ALA A 329	44.682	40.022	45.088	1.00	24.88
ATOM	2535	N	LEU A 330	42.290	40.354	47.161	1.00	25.47
ATOM	2536	CA	LEU A 330	41.672	41.133	48.219	1.00	26.16
ATOM	2537	C	LEU A 330	41.212	40.265	49.379	1.00	26.80
ATOM	2538	O	LEU A 330	40.994	40.761	50.471	1.00	27.00
ATOM	2539	CB	LEU A 330	40.504	41.930	47.669	1.00	26.08
ATOM	2540	CG	LEU A 330	40.954	42.981	46.653	1.00	26.04
ATOM	2541	CD1	LEU A 330	39.760	43.489	45.888	1.00	26.16
ATOM	2542	CD2	LEU A 330	41.688	44.134	47.353	1.00	26.65
ATOM	2543	N	GLY A 331	41.037	38.977	49.126	1.00	27.87
ATOM	2544	CA	GLY A 331	40.645	38.039	50.158	1.00	28.89
ATOM	2545	C	GLY A 331	39.176	38.086	50.526	1.00	29.43
ATOM	2546	O	GLY A 331	38.763	37.435	51.478	1.00	30.62
ATOM	2547	N	ASN A 332	38.400	38.887	49.808	1.00	29.59
ATOM	2548	CA	ASN A 332	36.963	38.995	50.023	1.00	29.64
ATOM	2549	C	ASN A 332	36.367	39.500	48.718	1.00	29.00
ATOM	2550	O	ASN A 332	36.649	40.605	48.303	1.00	28.70
ATOM	2551	CB	ASN A 332	36.656	39.973	51.156	1.00	30.15
ATOM	2552	CG	ASN A 332	35.162	40.094	51.441	1.00	31.82
ATOM	2553	OD1	ASN A 332	34.335	39.531	50.734	1.00	34.71
ATOM	2554	ND2	ASN A 332	34.818	40.818	52.504	1.00	35.11
ATOM	2555	N	PRO A 333	35.528	38.706	48.081	1.00	28.53
ATOM	2556	CA	PRO A 333	35.001	39.075	46.771	1.00	28.25
ATOM	2557	C	PRO A 333	34.176	40.343	46.807	1.00	27.91
ATOM	2558	O	PRO A 333	34.033	40.984	45.776	1.00	26.99
ATOM	2559	CB	PRO A 333	34.120	37.893	46.368	1.00	28.16
ATOM	2560	CG	PRO A 333	34.137	36.938	47.468	1.00	28.91
ATOM	2561	CD	PRO A 333	35.022	37.415	48.561	1.00	28.96
ATOM	2562	N	GLN A 334	33.638	40.701	47.965	1.00	27.80
ATOM	2563	CA	GLN A 334	32.814	41.896	48.039	1.00	27.97
ATOM	2564	C	GLN A 334	33.665	43.155	47.975	1.00	26.77
ATOM	2565	O	GLN A 334	33.144	44.245	47.764	1.00	26.51
ATOM	2566	CB	GLN A 334	31.906	41.867	49.278	1.00	28.98
ATOM	2567	CG	GLN A 334	30.675	40.969	49.038	1.00	32.17
ATOM	2568	CD	GLN A 334	29.661	40.962	50.175	1.00	35.54
ATOM	2569	OE1	GLN A 334	29.682	41.840	51.048	1.00	37.92
ATOM	2570	NE2	GLN A 334	28.760	39.961	50.163	1.00	36.72
ATOM	2571	N	GLU A 335	34.974	43.007	48.132	1.00	25.19
ATOM	2572	CA	GLU A 335	35.860	44.151	48.018	1.00	24.41
ATOM	2573	C	GLU A 335	36.170	44.466	46.537	1.00	22.68

C
C
C
C
N
C
C
O
N
C
C
O
C
C
C
O
N
C
C
O
C
C
C
O
N
C
C
C
O
N
C
C
C
O
N
C
C
C
O
N
C
C

ATOM	2574	O	GLU	A	335	36.700	45.521	46.228	1.00	21.42
ATOM	2575	CB	GLU	A	335	37.150	43.928	48.835	1.00	24.98
ATOM	2576	CG	GLU	A	335	36.974	44.148	50.343	1.00	27.94
ATOM	2577	CD	GLU	A	335	38.264	44.055	51.139	1.00	31.85
ATOM	2578	OE1	GLU	A	335	39.248	44.736	50.777	1.00	34.79
ATOM	2579	OE2	GLU	A	335	38.298	43.312	52.158	1.00	36.87
ATOM	2580	N	VAL	A	336	35.809	43.566	45.625	1.00	21.32
ATOM	2581	CA	VAL	A	336	36.113	43.751	44.201	1.00	20.37
ATOM	2582	C	VAL	A	336	35.541	45.034	43.603	1.00	20.01
ATOM	2583	O	VAL	A	336	36.247	45.798	42.954	1.00	19.07
ATOM	2584	CB	VAL	A	336	35.647	42.554	43.371	1.00	20.60
ATOM	2585	CG1	VAL	A	336	35.785	42.841	41.883	1.00	21.06
ATOM	2586	CG2	VAL	A	336	36.463	41.316	43.743	1.00	20.68
ATOM	2587	N	GLY	A	337	34.260	45.275	43.842	1.00	19.61
ATOM	2588	CA	GLY	A	337	33.593	46.437	43.317	1.00	19.45
ATOM	2589	C	GLY	A	337	34.205	47.760	43.731	1.00	19.51
ATOM	2590	O	GLY	A	337	34.522	48.576	42.871	1.00	18.93
ATOM	2591	N	PRO	A	338	34.333	48.010	45.032	1.00	20.41
ATOM	2592	CA	PRO	A	338	34.959	49.253	45.503	1.00	20.40
ATOM	2593	C	PRO	A	338	36.368	49.500	44.923	1.00	20.13
ATOM	2594	O	PRO	A	338	36.674	50.630	44.548	1.00	19.20
ATOM	2595	CB	PRO	A	338	34.960	49.092	47.035	1.00	20.63
ATOM	2596	CG	PRO	A	338	33.749	48.261	47.298	1.00	21.17
ATOM	2597	CD	PRO	A	338	33.782	47.221	46.151	1.00	20.95
ATOM	2598	N	LEU	A	339	37.199	48.470	44.831	1.00	20.34
ATOM	2599	CA	LEU	A	339	38.518	48.655	44.234	1.00	20.55
ATOM	2600	C	LEU	A	339	38.382	49.060	42.768	1.00	20.06
ATOM	2601	O	LEU	A	339	39.020	50.019	42.330	1.00	19.61
ATOM	2602	CB	LEU	A	339	39.383	47.408	44.358	1.00	20.81
ATOM	2603	CG	LEU	A	339	40.855	47.618	43.943	1.00	22.54
ATOM	2604	CD1	LEU	A	339	41.809	46.817	44.786	1.00	25.30
ATOM	2605	CD2	LEU	A	339	41.035	47.209	42.496	1.00	23.17
ATOM	2606	N	LEU	A	340	37.526	48.361	42.023	1.00	19.51
ATOM	2607	CA	LEU	A	340	37.327	48.687	40.600	1.00	19.36
ATOM	2608	C	LEU	A	340	36.827	50.120	40.427	1.00	19.30
ATOM	2609	O	LEU	A	340	37.318	50.861	39.566	1.00	17.55
ATOM	2610	CB	LEU	A	340	36.361	47.721	39.952	1.00	19.31
ATOM	2611	CG	LEU	A	340	36.929	46.333	39.637	1.00	21.99
ATOM	2612	CD1	LEU	A	340	35.842	45.506	39.069	1.00	22.03
ATOM	2613	CD2	LEU	A	340	38.140	46.390	38.673	1.00	21.20
ATOM	2614	N	ASN	A	341	35.879	50.521	41.270	1.00	19.17
ATOM	2615	CA	ASN	A	341	35.369	51.887	41.232	1.00	20.20
ATOM	2616	C	ASN	A	341	36.465	52.914	41.530	1.00	19.98
ATOM	2617	O	ASN	A	341	36.598	53.920	40.848	1.00	19.61
ATOM	2618	CB	ASN	A	341	34.181	52.043	42.196	1.00	20.66
ATOM	2619	CG	ASN	A	341	32.898	51.459	41.619	1.00	24.17
ATOM	2620	OD1	ASN	A	341	32.484	51.843	40.531	1.00	30.93
ATOM	2621	ND2	ASN	A	341	32.294	50.499	42.317	1.00	26.37
ATOM	2622	N	THR	A	342	37.245	52.633	42.558	1.00	20.07
ATOM	2623	CA	THR	A	342	38.351	53.489	42.939	1.00	20.49
ATOM	2624	C	THR	A	342	39.322	53.613	41.763	1.00	20.27
ATOM	2625	O	THR	A	342	39.835	54.688	41.502	1.00	19.66
ATOM	2626	CB	THR	A	342	39.001	52.909	44.189	1.00	20.82
ATOM	2627	OG1	THR	A	342	38.123	53.124	45.320	1.00	22.13
ATOM	2628	CG2	THR	A	342	40.308	53.625	44.545	1.00	21.01
ATOM	2629	N	MET	A	343	39.517	52.525	41.024	1.00	20.30
ATOM	2630	CA	MET	A	343	40.402	52.543	39.873	1.00	20.66
ATOM	2631	C	MET	A	343	39.932	53.445	38.758	1.00	20.29
ATOM	2632	O	MET	A	343	40.750	54.119	38.129	1.00	19.31
ATOM	2633	CB	MET	A	343	40.560	51.157	39.270	1.00	20.76
ATOM	2634	CG	MET	A	343	41.810	50.453	39.625	1.00	23.39

[illegible]

ATOM	2635	SD	MET	A	343	42.247	49.059	38.524	1.00	25.69
ATOM	2636	CE	MET	A	343	41.161	48.058	38.968	1.00	26.74
ATOM	2637	N	ILE	A	344	38.631	53.438	38.466	1.00	20.76
ATOM	2638	CA	ILE	A	344	38.167	54.187	37.312	1.00	21.23
ATOM	2639	C	ILE	A	344	37.648	55.577	37.565	1.00	20.71
ATOM	2640	O	ILE	A	344	37.763	56.403	36.666	1.00	20.65
ATOM	2641	CB	ILE	A	344	37.109	53.400	36.458	1.00	22.07
ATOM	2642	CG1	ILE	A	344	35.710	53.613	36.992	1.00	23.91
ATOM	2643	CG2	ILE	A	344	37.482	51.943	36.352	1.00	24.35
ATOM	2644	CD1	ILE	A	344	34.650	53.199	36.033	1.00	28.26
ATOM	2645	N	LYS	A	345	37.069	55.886	38.726	1.00	20.84
ATOM	2646	CA	LYS	A	345	36.482	57.231	38.817	1.00	21.15
ATOM	2647	C	LYS	A	345	37.464	58.376	38.784	1.00	19.86
ATOM	2648	O	LYS	A	345	38.459	58.429	39.517	1.00	19.23
ATOM	2649	CB	LYS	A	345	35.438	57.451	39.922	1.00	22.66
ATOM	2650	CG	LYS	A	345	35.562	56.714	41.190	1.00	27.82
ATOM	2651	CD	LYS	A	345	34.214	56.021	41.489	1.00	30.51
ATOM	2652	CE	LYS	A	345	33.398	56.782	42.502	1.00	32.51
ATOM	2653	NZ	LYS	A	345	34.069	56.806	43.829	1.00	37.63
ATOM	2654	N	GLY	A	346	37.143	59.307	37.899	1.00	18.18
ATOM	2655	CA	GLY	A	346	37.978	60.453	37.644	1.00	17.22
ATOM	2656	C	GLY	A	346	39.303	60.105	36.978	1.00	16.31
ATOM	2657	O	GLY	A	346	40.172	60.949	36.889	1.00	16.63
ATOM	2658	N	ARG	A	347	39.453	58.872	36.531	1.00	16.85
ATOM	2659	CA	ARG	A	347	40.697	58.413	35.904	1.00	17.52
ATOM	2660	C	ARG	A	347	40.469	57.811	34.502	1.00	17.57
ATOM	2661	O	ARG	A	347	41.178	58.143	33.559	1.00	17.10
ATOM	2662	CB	ARG	A	347	41.379	57.384	36.812	1.00	17.20
ATOM	2663	CG	ARG	A	347	41.822	57.949	38.181	1.00	16.77
ATOM	2664	CD	ARG	A	347	43.287	57.491	38.583	1.00	18.88
ATOM	2665	NE	ARG	A	347	43.254	56.087	38.618	1.00	17.77
ATOM	2666	CZ	ARG	A	347	44.115	55.220	38.149	1.00	16.11
ATOM	2667	NH1	ARG	A	347	45.323	55.512	37.658	1.00	16.59
ATOM	2668	NH2	ARG	A	347	43.734	53.978	38.276	1.00	13.22
ATOM	2669	N	TYR	A	348	39.472	56.954	34.375	1.00	18.74
ATOM	2670	CA	TYR	A	348	39.138	56.317	33.091	1.00	20.48
ATOM	2671	C	TYR	A	348	37.674	56.499	32.671	1.00	21.75
ATOM	2672	O	TYR	A	348	37.296	56.047	31.596	1.00	22.39
ATOM	2673	CB	TYR	A	348	39.411	54.800	33.124	1.00	19.97
ATOM	2674	CG	TYR	A	348	40.874	54.384	33.199	1.00	19.65
ATOM	2675	CD1	TYR	A	348	41.661	54.310	32.054	1.00	18.48
ATOM	2676	CD2	TYR	A	348	41.458	54.051	34.414	1.00	17.73
ATOM	2677	CE1	TYR	A	348	42.986	53.921	32.122	1.00	19.00
ATOM	2678	CE2	TYR	A	348	42.775	53.667	34.494	1.00	18.72
ATOM	2679	CZ	TYR	A	348	43.543	53.600	33.339	1.00	18.96
ATOM	2680	OH	TYR	A	348	44.856	53.203	33.419	1.00	17.77
ATOM	2681	N	ASN	A	349	36.837	57.123	33.488	1.00	23.99
ATOM	2682	CA	ASN	A	349	35.429	57.244	33.089	1.00	26.57
ATOM	2683	C	ASN	A	349	34.947	58.611	32.689	1.00	27.62
ATOM	2684	O	ASN	A	349	35.646	59.606	32.655	1.00	28.49
ATOM	2685	CB	ASN	A	349	34.496	56.720	34.150	1.00	26.11
ATOM	2686	CG	ASN	A	349	34.511	57.552	35.386	1.00	28.01
ATOM	2687	OD1	ASN	A	349	35.282	58.518	35.517	1.00	29.49
ATOM	2688	ND2	ASN	A	349	33.658	57.173	36.342	1.00	31.84
ATOM	2689	OXT	ASN	A	349	33.761	58.686	32.399	1.00	31.04
TER	2690		ASN	A	349					
ATOM	2691	N	LEU	S	795	45.837	35.555	30.600	1.00	35.49
ATOM	2692	CA	LEU	S	795	44.757	36.539	30.946	1.00	35.77
ATOM	2693	C	LEU	S	795	43.580	36.250	30.030	1.00	35.54
ATOM	2694	O	LEU	S	795	42.418	36.338	30.412	1.00	34.79
ATOM	2695	CB	LEU	S	795	45.257	37.967	30.787	1.00	36.01

[illegible]

ATOM	2696	CG	LEU	S	795	44.695	38.978	31.791	1.00	37.16
ATOM	2697	CD1	LEU	S	795	44.761	38.459	33.204	1.00	37.65
ATOM	2698	CD2	LEU	S	795	45.450	40.289	31.718	1.00	38.07
ATOM	2699	N	THR	S	796	43.936	35.940	28.796	1.00	35.47
ATOM	2700	CA	THR	S	796	43.060	35.351	27.800	1.00	36.25
ATOM	2701	C	THR	S	796	42.644	33.888	28.079	1.00	35.84
ATOM	2702	O	THR	S	796	41.819	33.331	27.365	1.00	36.38
ATOM	2703	CB	THR	S	796	43.817	35.429	26.457	1.00	36.63
ATOM	2704	OG1	THR	S	796	43.251	34.528	25.524	1.00	38.48
ATOM	2705	CG2	THR	S	796	45.257	34.907	26.593	1.00	37.07
ATOM	2706	N	SER	S	797	43.197	33.251	29.101	1.00	35.44
ATOM	2707	CA	SER	S	797	42.835	31.860	29.377	1.00	35.29
ATOM	2708	C	SER	S	797	41.426	31.760	29.984	1.00	35.18
ATOM	2709	O	SER	S	797	40.925	32.701	30.611	1.00	33.96
ATOM	2710	CB	SER	S	797	43.856	31.189	30.291	1.00	35.17
ATOM	2711	OG	SER	S	797	43.716	31.654	31.624	1.00	36.65
ATOM	2712	N	TYR	S	798	40.789	30.611	29.790	1.00	35.36
ATOM	2713	CA	TYR	S	798	39.427	30.430	30.256	1.00	35.94
ATOM	2714	C	TYR	S	798	39.148	29.075	30.890	1.00	35.14
ATOM	2715	O	TYR	S	798	39.845	28.095	30.657	1.00	35.54
ATOM	2716	CB	TYR	S	798	38.440	30.707	29.120	1.00	36.37
ATOM	2717	CG	TYR	S	798	38.554	29.794	27.920	1.00	39.89
ATOM	2718	CD1	TYR	S	798	39.587	29.934	26.996	1.00	42.41
ATOM	2719	CD2	TYR	S	798	37.606	28.808	27.697	1.00	43.13
ATOM	2720	CE1	TYR	S	798	39.681	29.095	25.893	1.00	43.97
ATOM	2721	CE2	TYR	S	798	37.687	27.963	26.597	1.00	44.94
ATOM	2722	CZ	TYR	S	798	38.723	28.109	25.698	1.00	44.95
ATOM	2723	OH	TYR	S	798	38.781	27.270	24.613	1.00	44.37
ATOM	2724	N	ASP	S	799	38.108	29.046	31.709	1.00	34.54
ATOM	2725	CA	ASP	S	799	37.685	27.840	32.392	1.00	33.87
ATOM	2726	C	ASP	S	799	36.600	27.187	31.545	1.00	32.58
ATOM	2727	O	ASP	S	799	36.465	27.501	30.366	1.00	32.32
ATOM	2728	CB	ASP	S	799	37.140	28.208	33.770	1.00	34.50
ATOM	2729	CG	ASP	S	799	37.299	27.104	34.773	1.00	36.07
ATOM	2730	OD1	ASP	S	799	36.790	25.986	34.551	1.00	37.15
ATOM	2731	OD2	ASP	S	799	37.918	27.279	35.833	1.00	41.54
ATOM	2732	N	CYS	S	800	35.812	26.301	32.141	1.00	30.83
ATOM	2733	CA	CYS	S	800	34.798	25.586	31.392	1.00	29.83
ATOM	2734	C	CYS	S	800	33.490	25.511	32.161	1.00	29.15
ATOM	2735	O	CYS	S	800	32.794	24.502	32.092	1.00	28.68
ATOM	2736	CB	CYS	S	800	35.273	24.167	31.092	1.00	29.46
ATOM	2737	SG	CYS	S	800	35.576	23.194	32.601	1.00	30.36
ATOM	2738	N	GLU	S	801	33.152	26.578	32.874	1.00	28.36
ATOM	2739	CA	GLU	S	801	31.936	26.605	33.670	1.00	28.63
ATOM	2740	C	GLU	S	801	30.667	26.793	32.829	1.00	28.34
ATOM	2741	O	GLU	S	801	30.626	27.562	31.867	1.00	27.30
ATOM	2742	CB	GLU	S	801	32.038	27.688	34.751	1.00	28.61
ATOM	2743	CG	GLU	S	801	33.252	27.501	35.655	1.00	30.83
ATOM	2744	CD	GLU	S	801	33.188	28.346	36.907	1.00	32.58
ATOM	2745	OE1	GLU	S	801	32.414	27.985	37.805	1.00	33.57
ATOM	2746	OE2	GLU	S	801	33.910	29.362	36.989	1.00	33.97
ATOM	2747	N	VAL	S	802	29.627	26.062	33.212	1.00	28.85
ATOM	2748	CA	VAL	S	802	28.350	26.079	32.515	1.00	29.03
ATOM	2749	C	VAL	S	802	27.233	26.056	33.546	1.00	29.80
ATOM	2750	O	VAL	S	802	27.505	25.948	34.726	1.00	29.92
ATOM	2751	CB	VAL	S	802	28.221	24.834	31.606	1.00	28.44
ATOM	2752	CG1	VAL	S	802	29.288	24.856	30.523	1.00	26.99
ATOM	2753	CG2	VAL	S	802	28.333	23.551	32.428	1.00	29.08
ATOM	2754	N	ASN	S	803	25.978	26.135	33.111	1.00	31.09
ATOM	2755	CA	ASN	S	803	24.853	26.053	34.042	1.00	32.53
ATOM	2756	C	ASN	S	803	24.550	24.641	34.497	1.00	33.81

ATOM	2757	O	ASN	S	803	23.456	24.143	34.270	1.00	34.27
ATOM	2758	CB	ASN	S	803	23.575	26.659	33.463	1.00	32.29
ATOM	2759	CG	ASN	S	803	23.640	28.146	33.367	1.00	31.77
ATOM	2760	OD1	ASN	S	803	24.688	28.741	33.616	1.00	33.47
ATOM	2761	ND2	ASN	S	803	22.525	28.772	33.005	1.00	29.83
ATOM	2762	N	ALA	S	804	25.521	24.015	35.147	1.00	35.23
ATOM	2763	CA	ALA	S	804	25.365	22.689	35.740	1.00	36.86
ATOM	2764	C	ALA	S	804	26.577	22.452	36.638	1.00	37.98
ATOM	2765	O	ALA	S	804	27.660	22.949	36.360	1.00	37.44
ATOM	2766	CB	ALA	S	804	25.285	21.610	34.676	1.00	36.63
ATOM	2767	N	PRO	S	805	26.394	21.694	37.711	1.00	40.20
ATOM	2768	CA	PRO	S	805	27.495	21.381	38.635	1.00	41.46
ATOM	2769	C	PRO	S	805	28.572	20.511	37.983	1.00	42.51
ATOM	2770	O	PRO	S	805	28.342	19.931	36.938	1.00	43.07
ATOM	2771	CB	PRO	S	805	26.799	20.615	39.774	1.00	41.54
ATOM	2772	CG	PRO	S	805	25.506	20.116	39.185	1.00	41.29
ATOM	2773	CD	PRO	S	805	25.117	21.076	38.115	1.00	40.40
ATOM	2774	N	ILE	S	806	29.728	20.406	38.622	1.00	44.25
ATOM	2775	CA	ILE	S	806	30.854	19.627	38.099	1.00	44.96
ATOM	2776	C	ILE	S	806	30.770	18.169	38.532	1.00	45.18
ATOM	2777	O	ILE	S	806	29.902	17.801	39.323	1.00	45.78
ATOM	2778	CB	ILE	S	806	32.197	20.246	38.569	1.00	45.19
ATOM	2779	CG1	ILE	S	806	32.412	20.018	40.070	1.00	46.10
ATOM	2780	CG2	ILE	S	806	32.230	21.743	38.246	1.00	46.08
ATOM	2781	CD1	ILE	S	806	33.740	20.574	40.597	1.00	46.98
ATOM	2782	N	LEU	S	812	29.934	8.629	39.561	1.00	43.80
ATOM	2783	CA	LEU	S	812	29.027	8.736	38.425	1.00	43.90
ATOM	2784	C	LEU	S	812	29.761	9.243	37.182	1.00	43.45
ATOM	2785	O	LEU	S	812	30.160	10.410	37.114	1.00	43.36
ATOM	2786	CB	LEU	S	812	27.862	9.678	38.757	1.00	44.22
ATOM	2787	CG	LEU	S	812	26.979	9.292	39.951	1.00	45.50
ATOM	2788	CD1	LEU	S	812	25.871	10.341	40.144	1.00	46.20
ATOM	2789	CD2	LEU	S	812	26.385	7.894	39.793	1.00	45.08
ATOM	2790	N	LEU	S	813	29.928	8.375	36.190	1.00	42.70
ATOM	2791	CA	LEU	S	813	30.620	8.776	34.969	1.00	42.21
ATOM	2792	C	LEU	S	813	29.711	9.604	34.057	1.00	41.46
ATOM	2793	O	LEU	S	813	28.492	9.438	34.062	1.00	41.23
ATOM	2794	CB	LEU	S	813	31.167	7.554	34.225	1.00	42.30
ATOM	2795	CG	LEU	S	813	32.093	6.644	35.046	1.00	42.35
ATOM	2796	CD1	LEU	S	813	32.494	5.419	34.233	1.00	42.48
ATOM	2797	CD2	LEU	S	813	33.322	7.398	35.527	1.00	41.80
ATOM	2798	N	GLN	S	814	30.326	10.492	33.283	1.00	40.74
ATOM	2799	CA	GLN	S	814	29.603	11.385	32.378	1.00	40.26
ATOM	2800	C	GLN	S	814	30.475	11.775	31.190	1.00	39.82
ATOM	2801	O	GLN	S	814	31.674	11.514	31.176	1.00	39.19
ATOM	2802	CB	GLN	S	814	29.173	12.649	33.122	1.00	40.24
ATOM	2803	CG	GLN	S	814	30.336	13.501	33.615	1.00	40.51
ATOM	2804	CD	GLN	S	814	29.879	14.725	34.406	1.00	41.88
ATOM	2805	OE1	GLN	S	814	29.200	14.590	35.419	1.00	41.86
ATOM	2806	NE2	GLN	S	814	30.253	15.919	33.940	1.00	40.63
ATOM	2807	N	GLY	S	815	29.864	12.416	30.200	1.00	39.81
ATOM	2808	CA	GLY	S	815	30.568	12.851	29.012	1.00	39.53
ATOM	2809	C	GLY	S	815	31.402	11.755	28.365	1.00	39.85
ATOM	2810	O	GLY	S	815	30.962	10.609	28.210	1.00	38.55
ATOM	2811	N	GLU	S	816	32.624	12.123	27.995	1.00	40.44
ATOM	2812	CA	GLU	S	816	33.553	11.208	27.352	1.00	41.24
ATOM	2813	C	GLU	S	816	33.744	9.926	28.154	1.00	41.73
ATOM	2814	O	GLU	S	816	33.895	8.852	27.577	1.00	41.11
ATOM	2815	CB	GLU	S	816	34.909	11.884	27.148	1.00	41.24
ATOM	2816	CG	GLU	S	816	35.752	11.238	26.063	1.00	42.19
ATOM	2817	CD	GLU	S	816	37.161	11.790	26.019	1.00	43.70

O
C
C
O
N
N
C
C
O
C
N
C
C
O
C
C
N
C
C
O
C
C
C
N
C
C
O
C
C
C
C
N
C
C
O
C
C
C
C
O
N
N
C
C
O
N
C
C
O
C
C
C

ATOM	2818	OE1	GLU	S	816	37.985	11.311	26.814	1.00	45.25
ATOM	2819	OE2	GLU	S	816	37.447	12.696	25.201	1.00	44.50
ATOM	2820	N	GLU	S	817	33.724	10.039	29.479	1.00	42.66
ATOM	2821	CA	GLU	S	817	33.925	8.877	30.340	1.00	43.73
ATOM	2822	C	GLU	S	817	32.744	7.917	30.268	1.00	44.19
ATOM	2823	O	GLU	S	817	32.930	6.699	30.252	1.00	43.99
ATOM	2824	CB	GLU	S	817	34.167	9.308	31.788	1.00	43.81
ATOM	2825	CG	GLU	S	817	35.463	10.077	31.989	1.00	44.98
ATOM	2826	CD	GLU	S	817	35.337	11.569	31.688	1.00	46.95
ATOM	2827	OE1	GLU	S	817	34.221	12.052	31.394	1.00	48.87
ATOM	2828	OE2	GLU	S	817	36.362	12.277	31.748	1.00	49.02
ATOM	2829	N	LEU	S	818	31.536	8.470	30.231	1.00	44.93
ATOM	2830	CA	LEU	S	818	30.335	7.663	30.149	1.00	45.78
ATOM	2831	C	LEU	S	818	30.359	6.882	28.846	1.00	46.89
ATOM	2832	O	LEU	S	818	30.163	5.663	28.833	1.00	46.50
ATOM	2833	CB	LEU	S	818	29.077	8.537	30.219	1.00	45.65
ATOM	2834	CG	LEU	S	818	27.741	7.788	30.147	1.00	45.58
ATOM	2835	CD1	LEU	S	818	27.612	6.809	31.318	1.00	45.36
ATOM	2836	CD2	LEU	S	818	26.551	8.719	30.133	1.00	43.95
ATOM	2837	N	LEU	S	819	30.654	7.588	27.760	1.00	48.01
ATOM	2838	CA	LEU	S	819	30.607	7.014	26.420	1.00	49.24
ATOM	2839	C	LEU	S	819	31.559	5.829	26.247	1.00	50.32
ATOM	2840	O	LEU	S	819	31.169	4.777	25.731	1.00	50.11
ATOM	2841	CB	LEU	S	819	30.903	8.105	25.380	1.00	49.22
ATOM	2842	CG	LEU	S	819	30.848	7.724	23.901	1.00	49.32
ATOM	2843	CD1	LEU	S	819	29.495	7.178	23.497	1.00	48.73
ATOM	2844	CD2	LEU	S	819	31.205	8.940	23.058	1.00	49.85
ATOM	2845	N	ARG	S	820	32.801	6.005	26.684	1.00	51.33
ATOM	2846	CA	ARG	S	820	33.807	4.966	26.550	1.00	52.53
ATOM	2847	C	ARG	S	820	33.471	3.748	27.414	1.00	52.95
ATOM	2848	O	ARG	S	820	33.534	2.606	26.942	1.00	53.06
ATOM	2849	CB	ARG	S	820	35.185	5.529	26.898	1.00	52.91
ATOM	2850	CG	ARG	S	820	35.620	6.583	25.904	1.00	54.20
ATOM	2851	CD	ARG	S	820	37.044	7.046	26.040	1.00	55.94
ATOM	2852	NE	ARG	S	820	37.320	8.113	25.081	1.00	58.09
ATOM	2853	CZ	ARG	S	820	38.453	8.808	25.022	1.00	59.82
ATOM	2854	NH1	ARG	S	820	39.448	8.555	25.867	1.00	60.28
ATOM	2855	NH2	ARG	S	820	38.590	9.765	24.108	1.00	60.61
ATOM	2856	N	ALA	S	821	33.106	3.994	28.670	1.00	53.36
ATOM	2857	CA	ALA	S	821	32.698	2.920	29.561	1.00	53.67
ATOM	2858	C	ALA	S	821	31.598	2.094	28.892	1.00	54.08
ATOM	2859	O	ALA	S	821	31.648	0.862	28.897	1.00	54.01
ATOM	2860	CB	ALA	S	821	32.215	3.479	30.883	1.00	53.55
ATOM	2861	N	LEU	S	822	30.616	2.777	28.308	1.00	54.44
ATOM	2862	CA	LEU	S	822	29.516	2.110	27.622	1.00	54.93
ATOM	2863	C	LEU	S	822	30.007	1.342	26.398	1.00	55.41
ATOM	2864	O	LEU	S	822	29.557	0.230	26.143	1.00	55.60
ATOM	2865	CB	LEU	S	822	28.445	3.117	27.212	1.00	54.90
ATOM	2866	CG	LEU	S	822	27.667	3.756	28.361	1.00	54.80
ATOM	2867	CD1	LEU	S	822	26.531	4.601	27.801	1.00	54.39
ATOM	2868	CD2	LEU	S	822	27.139	2.702	29.335	1.00	54.48
ATOM	2869	N	ASP	S	823	30.911	1.949	25.634	1.00	55.82
ATOM	2870	CA	ASP	S	823	31.525	1.282	24.492	1.00	56.04
ATOM	2871	C	ASP	S	823	32.893	0.735	24.901	1.00	56.10
ATOM	2872	O	ASP	S	823	33.186	-0.451	24.731	1.00	56.09
ATOM	2873	CB	ASP	S	823	31.688	2.252	23.320	1.00	56.21
ATOM	2874	CG	ASP	S	823	31.858	1.535	21.997	1.00	56.45
ATOM	2875	OD1	ASP	S	823	32.245	0.351	22.028	1.00	58.65
ATOM	2876	OD2	ASP	S	823	31.628	2.056	20.885	1.00	56.19
TER	2877		ASP	S	823					
HETATM	2878	FE	FE2	Al350		23.294	27.501	28.594	1.00	20.46

[illegible]**FE**

HETATM	2879	C1	OGA	A1351	22.091	25.173	27.594	1.00	24.79	
HETATM	2880	C2	OGA	A1351	21.066	25.829	28.202	1.00	24.27	
HETATM	2881	C4	OGA	A1351	18.756	25.714	29.004	1.00	23.09	
HETATM	2882	C5	OGA	A1351	17.415	25.241	28.495	1.00	23.17	
HETATM	2883	O1	OGA	A1351	21.909	24.061	27.090	1.00	25.24	
HETATM	2884	O2	OGA	A1351	23.219	25.658	27.531	1.00	24.40	
HETATM	2885	O2	OGA	A1351	21.192	26.959	28.711	1.00	21.19	
HETATM	2886	O3	OGA	A1351	16.416	25.662	29.055	1.00	23.03	
HETATM	2887	N1	OGA	A1351	19.886	25.203	28.228	1.00	21.70	
HETATM	2888	O4	OGA	A1351	17.332	24.475	27.537	1.00	23.98	
HETATM	2889	S	SO4	A1352	0.316	25.182	43.602	1.00	77.77	
HETATM	2890	O1	SO4	A1352	1.239	25.980	44.403	1.00	77.64	
HETATM	2891	O2	SO4	A1352	1.075	24.260	42.760	1.00	77.88	
HETATM	2892	O3	SO4	A1352	-0.525	24.416	44.514	1.00	78.38	
HETATM	2893	O4	SO4	A1352	-0.507	26.042	42.757	1.00	76.90	
HETATM	2894	S	SO4	A1353	1.990	28.487	29.834	1.00	69.20	
HETATM	2895	O1	SO4	A1353	3.243	29.065	30.309	1.00	68.34	
HETATM	2896	O2	SO4	A1353	2.236	27.438	28.847	1.00	67.90	
HETATM	2897	O3	SO4	A1353	1.298	27.948	31.009	1.00	70.32	
HETATM	2898	O4	SO4	A1353	1.162	29.517	29.203	1.00	69.63	
HETATM	2899	O	HOH	H	1	35.955	31.618	40.285	1.00	80.01
HETATM	2900	O	HOH	H	2	38.513	33.804	31.613	1.00	33.04
HETATM	2901	O	HOH	H	3	36.648	25.786	38.779	1.00	76.96
HETATM	2902	O	HOH	H	4	38.106	25.337	29.179	1.00	54.79
HETATM	2903	O	HOH	H	5	34.990	30.561	34.967	1.00	30.13
HETATM	2904	O	HOH	H	6	33.934	31.237	38.711	1.00	40.66
HETATM	2905	O	HOH	H	7	30.766	25.787	37.613	1.00	54.75
HETATM	2906	O	HOH	H	8	33.667	28.867	40.196	1.00	59.66
HETATM	2907	O	HOH	H	9	28.622	27.043	37.556	1.00	57.58
HETATM	2908	O	HOH	H	10	19.894	26.655	33.706	1.00	54.88
HETATM	2909	O	HOH	H	11	30.052	24.213	35.628	1.00	41.23
HETATM	2910	O	HOH	H	12	28.737	12.960	37.083	1.00	59.80
HETATM	2911	O	HOH	H	13	35.568	13.822	23.888	1.00	38.00
HETATM	2912	O	HOH	H	14	30.722	-1.323	21.296	1.00	48.92
HETATM	2913	O	HOH	H	15	32.110	2.136	17.673	1.00	69.92
HETATM	2914	O	HOH	Z	1	9.466	21.720	12.039	1.00	63.79
HETATM	2915	O	HOH	Z	2	1.367	21.270	7.724	1.00	60.01
HETATM	2916	O	HOH	Z	3	3.426	13.325	8.811	1.00	43.04
HETATM	2917	O	HOH	Z	4	-0.760	13.029	7.574	1.00	47.08
HETATM	2918	O	HOH	Z	5	2.515	19.304	5.195	1.00	46.76
HETATM	2919	O	HOH	Z	6	4.861	33.534	13.331	1.00	75.60
HETATM	2920	O	HOH	Z	7	1.403	29.250	13.007	1.00	46.80
HETATM	2921	O	HOH	Z	8	1.614				

HETATM	2940	O	HOH	Z	27	23.279	21.788	0.672	1.00	62.23	O
HETATM	2941	O	HOH	Z	28	27.443	22.009	43.177	1.00	68.81	O
HETATM	2942	O	HOH	Z	29	27.326	30.900	5.769	1.00	84.31	O
HETATM	2943	O	HOH	Z	30	16.938	35.662	41.749	1.00	51.88	O
HETATM	2944	O	HOH	Z	31	36.792	29.262	21.033	1.00	42.38	O
HETATM	2945	O	HOH	Z	32	26.719	37.403	13.167	1.00	60.20	O
HETATM	2946	O	HOH	Z	33	29.797	37.021	10.379	1.00	60.24	O
HETATM	2947	O	HOH	Z	34	28.365	37.713	15.023	1.00	68.08	O
HETATM	2948	O	HOH	Z	35	27.471	34.815	9.298	1.00	63.90	O
HETATM	2949	O	HOH	Z	36	24.262	32.919	12.792	1.00	45.02	O
HETATM	2950	O	HOH	Z	37	19.704	17.909	13.178	1.00	28.78	O
HETATM	2951	O	HOH	Z	38	22.022	12.870	8.792	1.00	48.37	O
HETATM	2952	O	HOH	Z	39	18.151	14.971	12.982	1.00	36.76	O
HETATM	2953	O	HOH	Z	40	29.160	5.439	16.977	1.00	44.05	O
HETATM	2954	O	HOH	Z	41	18.863	14.590	16.204	1.00	30.84	O
HETATM	2955	O	HOH	Z	42	12.149	5.293	13.385	1.00	72.13	O
HETATM	2956	O	HOH	Z	43	15.651	2.782	13.845	1.00	41.52	O
HETATM	2957	O	HOH	Z	44	14.014	7.467	18.234	1.00	52.22	O
HETATM	2958	O	HOH	Z	45	5.548	12.548	27.846	1.00	38.03	O
HETATM	2959	O	HOH	Z	46	12.742	5.782	36.187	1.00	51.07	O
HETATM	2960	O	HOH	Z	47	19.063	6.567	36.600	1.00	49.16	O
HETATM	2961	O	HOH	Z	48	19.545	2.633	38.104	1.00	67.28	O
HETATM	2962	O	HOH	Z	49	7.710	14.276	19.473	1.00	48.41	O
HETATM	2963	O	HOH	Z	50	15.732	7.234	38.833	1.00	54.12	O
HETATM	2964	O	HOH	Z	51	21.932	13.291	44.351	1.00	62.49	O
HETATM	2965	O	HOH	Z	52	33.998	44.086	25.334	1.00	53.56	O
HETATM	2966	O	HOH	Z	53	12.673	21.178	43.612	1.00	63.01	O
HETATM	2967	O	HOH	Z	54	8.172	26.738	44.107	1.00	61.46	O
HETATM	2968	O	HOH	Z	55	9.613	30.854	42.520	1.00	54.56	O
HETATM	2969	O	HOH	Z	56	13.563	35.806	31.131	1.00	39.09	O
HETATM	2970	O	HOH	Z	57	15.688	37.473	35.304	1.00	47.58	O
HETATM	2971	O	HOH	Z	58	7.422	43.868	25.982	1.00	75.57	O
HETATM	2972	O	HOH	Z	59	7.978	38.223	34.865	1.00	59.51	O
HETATM	2973	O	HOH	Z	60	16.338	30.836	40.223	1.00	38.80	O
HETATM	2974	O	HOH	Z	61	17.035	27.760	38.288	1.00	52.22	O
HETATM	2975	O	HOH	Z	62	22.131	25.023	41.390	1.00	55.16	O
HETATM	2976	O	HOH	Z	63	29.869	29.910	39.122	1.00	47.28	O
HETATM	2977	O	HOH	Z	64	28.353	24.399	41.766	1.00	60.35	O
HETATM	2978	O	HOH	Z	65	31.794	27.570	41.962	1.00	48.59	O
HETATM	2979	O	HOH	Z	66	28.058	28.695	48.927	1.00	65.95	O
HETATM	2980	O	HOH	Z	67	24.838	24.783	42.190	1.00	52.90	O
HETATM	2981	O	HOH	Z	68	11.541	32.183	15.082	1.00	63.99	O
HETATM	2982	O	HOH	Z	69	31.599	33.767	45.823	1.00	44.28	O
HETATM	2983	O	HOH	Z	70	24.728	38.721	49.282	1.00	48.25	O
HETATM	2984	O	HOH	Z	71	16.271	36.399	44.087	1.00	58.46	O
HETATM	2985	O	HOH	Z	72	17.845	37.716	46.244	1.00	57.93	O
HETATM	2986	O	HOH	Z	73	16.480	33.117	41.520	1.00	59.15	O
HETATM	2987	O	HOH	Z	74	40.791	21.415	26.920	1.00	58.98	O
HETATM	2988	O	HOH	Z	75	21.842	17.819	48.106	1.00	67.11	O
HETATM	2989	O	HOH	Z	76	8.791	17.468	46.626	1.00	63.28	O
HETATM	2990	O	HOH	Z	77	17.141	16.914	47.607	1.00	52.87	O
HETATM	2991	O	HOH	Z	78	21.626	14.804	40.702	1.00	53.91	O
HETATM	2992	O	HOH	Z	79	39.117	50.091	47.735	1.00	57.17	O
HETATM	2993	O	HOH	Z	80	10.617	19.257	44.587	1.00	69.00	O
HETATM	2994	O	HOH	Z	81	1.682	24.435	36.842	1.00	55.40	O
HETATM	2995	O	HOH	Z	82	4.627	30.781	36.487	1.00	53.38	O
HETATM	2996	O	HOH	Z	83	17.463	26.906	33.818	1.00	38.64	O
HETATM	2997	O	HOH	Z	84	18.429	25.785	36.464	1.00	54.65	O
HETATM	2998	O	HOH	Z	85	23.466	17.336	36.578	1.00	41.94	O
HETATM	2999	O	HOH	Z	86	26.890	12.949	30.365	1.00	49.34	O
HETATM	3000	O	HOH	Z	87	21.694	10.405	34.333	1.00	45.95	O

HETATM	3001	O	HOH	Z	88	20.030	9.589	36.847	1.00	52.33
HETATM	3002	O	HOH	Z	89	18.447	-1.706	32.981	1.00	66.29
HETATM	3003	O	HOH	Z	90	16.300	-0.205	30.017	1.00	50.70
HETATM	3004	O	HOH	Z	91	17.950	1.645	20.589	1.00	51.55
HETATM	3005	O	HOH	Z	92	26.301	5.439	16.918	1.00	36.67
HETATM	3006	O	HOH	Z	93	33.944	10.218	13.383	1.00	51.60
HETATM	3007	O	HOH	Z	94	30.893	16.371	11.174	1.00	40.00
HETATM	3008	O	HOH	Z	95	32.606	13.689	20.709	1.00	45.01
HETATM	3009	O	HOH	Z	96	31.860	10.158	7.765	1.00	58.55
HETATM	3010	O	HOH	Z	97	36.957	10.422	7.614	1.00	76.91
HETATM	3011	O	HOH	Z	98	35.951	16.836	31.735	1.00	59.28
HETATM	3012	O	HOH	Z	99	39.867	18.412	27.150	1.00	50.38
HETATM	3013	O	HOH	Z	100	13.436	20.952	28.355	1.00	27.89
HETATM	3014	O	HOH	Z	101	3.992	21.265	30.540	1.00	43.76
HETATM	3015	O	HOH	Z	102	30.735	37.910	33.103	1.00	30.45
HETATM	3016	O	HOH	Z	103	25.986	26.303	26.047	1.00	26.08
HETATM	3017	O	HOH	Z	104	36.837	32.025	33.001	1.00	37.86
HETATM	3018	O	HOH	Z	105	35.845	25.360	27.653	1.00	33.53
HETATM	3019	O	HOH	Z	106	31.874	20.474	33.040	1.00	55.47
HETATM	3020	O	HOH	Z	107	36.793	26.619	20.918	1.00	34.57
HETATM	3021	O	HOH	Z	108	17.114	16.909	17.862	1.00	35.83
HETATM	3022	O	HOH	Z	109	9.968	13.510	19.464	1.00	37.10
HETATM	3023	O	HOH	Z	110	5.274	16.717	22.023	1.00	44.26
HETATM	3024	O	HOH	Z	111	7.041	16.862	20.149	1.00	37.71
HETATM	3025	O	HOH	Z	112	6.517	22.763	22.963	1.00	38.94
HETATM	3026	O	HOH	Z	113	29.471	38.812	26.249	1.00	22.19
HETATM	3027	O	HOH	Z	114	32.253	43.121	24.229	1.00	53.95
HETATM	3028	O	HOH	Z	115	28.797	40.227	16.136	1.00	64.15
HETATM	3029	O	HOH	Z	116	30.645	39.042	18.925	1.00	33.21
HETATM	3030	O	HOH	Z	117	25.466	42.302	17.883	1.00	64.00
HETATM	3031	O	HOH	Z	118	27.698	44.486	31.068	1.00	26.13
HETATM	3032	O	HOH	Z	119	30.274	44.879	25.031	1.00	38.32
HETATM	3033	O	HOH	Z	120	27.635	48.525	29.776	1.00	35.59
HETATM	3034	O	HOH	Z	121	25.933	50.527	30.463	1.00	35.79
HETATM	3035	O	HOH	Z	122	26.614	31.094	38.158	1.00	32.11
HETATM	3036	O	HOH	Z	123	29.486	36.934	35.226	1.00	27.40
HETATM	3037	O	HOH	Z	124	30.510	30.229	36.300	1.00	31.99
HETATM	3038	O	HOH	Z	125	19.859	31.575	33.786	1.00	31.77
HETATM	3039	O	HOH	Z	126	18.490	38.301	36.415	1.00	43.54
HETATM	3040	O	HOH	Z	127	19.538	36.155	33.217	1.00	30.91
HETATM	3041	O	HOH	Z	128	25.386	44.358	40.124	1.00	54.33
HETATM	3042	O	HOH	Z	129	21.227				

[illegible]

HETATM	3062	O	HOH	Z	149	14.521	18.471	12.156	1.00	37.16	
HETATM	3063	O	HOH	Z	150	33.566	26.082	16.801	1.00	38.34	
HETATM	3064	O	HOH	Z	151	27.798	27.494	24.390	1.00	25.11	
HETATM	3065	O	HOH	Z	152	30.253	36.176	26.688	1.00	27.79	
HETATM	3066	O	HOH	Z	153	32.762	34.069	19.065	1.00	47.61	
HETATM	3067	O	HOH	Z	154	27.363	30.094	25.612	1.00	23.74	
HETATM	3068	O	HOH	Z	155	16.282	36.168	29.566	1.00	33.13	
HETATM	3069	O	HOH	Z	156	13.289	26.522	28.472	1.00	47.17	
HETATM	3070	O	HOH	Z	157	7.556	30.888	24.716	1.00	44.35	
HETATM	3071	O	HOH	Z	158	7.790	30.392	27.265	1.00	37.71	
HETATM	3072	O	HOH	Z	159	3.910	21.360	21.198	1.00	60.23	
HETATM	3073	O	HOH	Z	160	6.411	23.837	25.422	1.00	40.51	
HETATM	3074	O	HOH	Z	161	3.001	17.561	27.759	1.00	55.61	
HETATM	3075	O	HOH	Z	162	5.352	18.754	24.064	1.00	42.17	
HETATM	3076	O	HOH	Z	163	40.897	21.985	24.633	1.00	54.83	
HETATM	3077	O	HOH	Z	164	42.626	37.977	17.732	1.00	71.69	
HETATM	3078	O	HOH	Z	165	42.463	44.114	19.559	1.00	54.59	
HETATM	3079	O	HOH	Z	166	33.754	36.855	19.984	1.00	36.17	
HETATM	3080	O	HOH	Z	167	33.194	39.271	33.406	1.00	28.26	
HETATM	3081	O	HOH	Z	168	37.482	29.751	37.743	1.00	57.74	
HETATM	3082	O	HOH	Z	169	38.789	34.566	42.752	1.00	40.46	
HETATM	3083	O	HOH	Z	170	32.020	40.390	44.008	1.00	38.46	
HETATM	3084	O	HOH	Z	171	31.865	44.151	45.174	1.00	45.64	
HETATM	3085	O	HOH	Z	172	41.427	43.475	51.164	1.00	50.64	
HETATM	3086	O	HOH	Z	173	38.498	47.344	48.085	1.00	60.19	
HETATM	3087	O	HOH	Z	174	35.239	52.784	45.826	1.00	54.63	
HETATM	3088	O	HOH	Z	175	32.343	51.933	45.286	1.00	58.03	
HETATM	3089	O	HOH	Z	176	39.267	57.239	41.745	1.00	26.45	
HETATM	3090	O	HOH	Z	177	32.755	59.287	44.804	1.00	60.46	
HETATM	3091	O	HOH	Z	178	46.839	53.182	31.757	1.00	31.38	
HETATM	3092	O	HOH	Z	179	37.840	55.111	29.241	1.00	44.20	
CONNECT	1478	2878									
CONNECT	1498	2878									
CONNECT	2167	2878									
CONNECT	2878	2885	2884	1478	2167	1498					
CONNECT	2879	2880	2883	2884							
CONNECT	2880	2879	2885	2887							
CONNECT	2881	2882	2887								
CONNECT	2882	2881	2886	2888							
CONNECT	2883	2879									
CONNECT	2884	2879	2878								
CONNECT	2885	2880	2878								
CONNECT	2886	2882									
CONNECT	2887	2880	2881								

Structure 2

Below are the coordinates for structure 2 (the 2.25 Å structure of FIH:Fe(II):2OG:CAD):

```

HEADER      TRANSCRIPTION ACTIVATOR/INHIBITOR          12-AUG-02   1H2L
TITLE       FACTOR INHIBITING HIF-1 ALPHA IN COMPLEX WITH HIF-1 ALPHA
TITLE       2 FRAGMENT PEPTIDE
COMPND      MOL_ID: 1;
COMPND      2 MOLECULE: FACTOR INHIBITING HIF1;
COMPND      3 CHAIN: A;
COMPND      4 ENGINEERED: YES;
COMPND      5 MOL_ID: 2;
COMPND      6 MOLECULE: HYPOXIA-INDUCIBLE FACTOR 1 ALPHA;
COMPND      7 SYNONYM: HIF-1 ALPHA, ARNT INTERACTING PROTEIN,
COMPND      8 MEMBER OF PAS PROTEIN 1;
COMPND      9 CHAIN: S;
COMPND     10 FRAGMENT: C-TERMINAL TRANSACTIVATION DOMAIN FRAGMENT
COMPND     11 RESIDUES 786-826
SOURCE      MOL_ID: 1;
SOURCE      2 ORGANISM_SCIENTIFIC: HOMO SAPIENS;
SOURCE      3 ORGANISM_COMMON: HUMAN;
SOURCE      4 EXPRESSION_SYSTEM: ESCHERICHIA COLI;
SOURCE      5 EXPRESSION_SYSTEM_STRAIN: BL21(DE3);
SOURCE      6 EXPRESSION_SYSTEM_PLASMID: PET28A(+);
SOURCE      7 MOL_ID: 2;
SOURCE      8 ORGANISM_SCIENTIFIC: HOMO SAPIENS;
SOURCE      9 ORGANISM_COMMON: HUMAN;
SOURCE     10 EXPRESSION_SYSTEM: ESCHERICHIA COLI;
SOURCE     11 EXPRESSION_SYSTEM_STRAIN: BL21(DE3);
SOURCE     12 EXPRESSION_SYSTEM_PLASMID: PGEX-GP-1
KEYWDS      FIH, HIF, DSBH, OXYGENASE, TRANSCRIPTION, HYPOXIA,
KEYWDS      2 2-OXOGLUTARATE, ASPARAGINYL HYDROXYLASE, HYDROXYLASE
EXPDTA      X-RAY DIFFRACTION
AUTHOR      J.M.ELKINS,K.S.HEWITSON,L.A.MCNEILL,I.SCHLEMMINGER,
AUTHOR      2 J.F.SEIBEL,C.J.SCHOFIELD
REVDAT      1 04-SEP-02 1H2L 0
JRNL        AUTH J.M.ELKINS,K.S.HEWITSON,L.A.MCNEILL,
JRNL        AUTH 2 I.SCHLEMMINGER,J.F.SEIBEL,C.J.SCHOFIELD
JRNL        TITL FIH:HIF-FRAGMENT COMPLEXES
JRNL        REF TO BE PUBLISHED
JRNL        REFN
REMARK      2
REMARK      2 RESOLUTION. 2.25 ANGSTROMS.
REMARK      3
REMARK      3 REFINEMENT.
REMARK      3 PROGRAM : REFMAC 5.0
REMARK      3 AUTHORS : MURSHUDOV,VAGIN,DODSON
REMARK      3
REMARK      3 REFINEMENT TARGET : MAXIMUM LIKELIHOOD
REMARK      3
REMARK      3 DATA USED IN REFINEMENT.
REMARK      3 RESOLUTION RANGE HIGH (ANGSTROMS) : 2.25
REMARK      3 RESOLUTION RANGE LOW (ANGSTROMS) : 38.00
REMARK      3 DATA CUTOFF (SIGMA(F)) : NONE
REMARK      3 COMPLETENESS FOR RANGE (%) : 99.98
REMARK      3 NUMBER OF REFLECTIONS : 25127
REMARK      3
REMARK      3 FIT TO DATA USED IN REFINEMENT.
REMARK      3 CROSS-VALIDATION METHOD : THROUGHOUT

```

REMARK 3 FREE R VALUE TEST SET SELECTION : RANDOM
 REMARK 3 R VALUE (WORKING + TEST SET) : 0.18514
 REMARK 3 R VALUE (WORKING SET) : 0.18253
 REMARK 3 FREE R VALUE : 0.21738
 REMARK 3 FREE R VALUE TEST SET SIZE (%) : 7.7
 REMARK 3 FREE R VALUE TEST SET COUNT : 2104
 REMARK 3
 REMARK 3 FIT IN THE HIGHEST RESOLUTION BIN.
 REMARK 3 TOTAL NUMBER OF BINS USED : 20
 REMARK 3 BIN RESOLUTION RANGE HIGH : 2.250
 REMARK 3 BIN RESOLUTION RANGE LOW : 2.308
 REMARK 3 REFLECTION IN BIN (WORKING SET) : 1783
 REMARK 3 BIN R VALUE (WORKING SET) : 0.194
 REMARK 3 BIN FREE R VALUE SET COUNT : 170
 REMARK 3 BIN FREE R VALUE : 0.228
 REMARK 3
 REMARK 3 NUMBER OF NON-HYDROGEN ATOMS USED IN REFINEMENT.
 REMARK 3 PROTEIN ATOMS : 2863
 REMARK 3 NUCLEIC ACID ATOMS : 0
 REMARK 3 HETEROGEN ATOMS : 21
 REMARK 3 SOLVENT ATOMS : 139
 REMARK 3
 REMARK 3 B VALUES.
 REMARK 3 FROM WILSON PLOT (A**2) : NULL
 REMARK 3 MEAN B VALUE (OVERALL, A**2) : 27.234
 REMARK 3 OVERALL ANISOTROPIC B VALUE.
 REMARK 3 B11 (A**2) : -0.40
 REMARK 3 B22 (A**2) : -0.40
 REMARK 3 B33 (A**2) : 0.80
 REMARK 3 B12 (A**2) : 0.00
 REMARK 3 B13 (A**2) : 0.00
 REMARK 3 B23 (A**2) : 0.00
 REMARK 3
 REMARK 3 ESTIMATED OVERALL COORDINATE ERROR.
 REMARK 3 ESU BASED ON R VALUE (A) : 0.203
 REMARK 3 ESU BASED ON FREE R VALUE (A) : 0.174
 REMARK 3 ESU BASED ON MAXIMUM LIKELIHOOD (A) : 0.165
 REMARK 3 ESU FOR B VALUES BASED ON MAXIMUM LIKELIHOOD (A**2) : 6.444
 REMARK 3
 REMARK 3 CORRELATION COEFFICIENTS.
 REMARK 3 CORRELATION COEFFICIENT FO-FC : 0.956
 REMARK 3 CORRELATION COEFFICIENT FO-FC FREE : 0.939
 REMARK 3
 REMARK 3 RMS DEVIATIONS FROM IDEAL VALUES COUNT RMS WEIGHT
 REMARK 3 BOND LENGTHS REFINED (A) : 2961 ; 0.013 ; 0.021
 REMARK 3 BOND LENGTHS REFINED (A) : 2961 ; 0.013 ; 0.021
 REMARK 3 BOND LENGTHS OTHERS (A) : 2554 ; 0.001 ; 0.020
 REMARK 3 BOND ANGLES REFINED (DEGREES) : 4026 ; 1.404 ; 1.949
 REMARK 3 BOND ANGLES OTHERS (DEGREES) : 5966 ; 0.727 ; 3.000
 REMARK 3 TORSION ANGLES, PERIOD 1 (DEGREES) : 350 ; 4.037 ; 3.000
 REMARK 3 TORSION ANGLES, PERIOD 3 (DEGREES) : 515 ; 18.189 ; 15.000
 REMARK 3 CHIRAL-CENTER RESTRAINTS (A**3) : 413 ; 0.085 ; 0.200
 REMARK 3 GENERAL PLANES REFINED (A) : 3315 ; 0.005 ; 0.020
 REMARK 3 GENERAL PLANES OTHERS (A) : 602 ; 0.002 ; 0.020
 REMARK 3 NON-BONDED CONTACTS REFINED (A) : 693 ; 0.221 ; 0.300
 REMARK 3 NON-BONDED CONTACTS OTHERS (A) : 2483 ; 0.204 ; 0.300
 REMARK 3 H-BOND (X...Y) REFINED (A) : 208 ; 0.156 ; 0.500
 REMARK 3 SYMMETRY VDW REFINED (A) : 16 ; 0.256 ; 0.300
 REMARK 3 SYMMETRY VDW OTHERS (A) : 63 ; 0.259 ; 0.300
 REMARK 3 SYMMETRY H-BOND REFINED (A) : 10 ; 0.200 ; 0.500
 REMARK 3

REMARK 3 ISOTROPIC THERMAL FACTOR RESTRAINTS. COUNT RMS WEIGHT
REMARK 3 MAIN-CHAIN BOND REFINED (A**2): 1767 ; 0.649 ; 1.500
REMARK 3 MAIN-CHAIN ANGLE REFINED (A**2): 2847 ; 1.227 ; 2.000
REMARK 3 SIDE-CHAIN BOND REFINED (A**2): 1194 ; 1.887 ; 3.000
REMARK 3 SIDE-CHAIN ANGLE REFINED (A**2): 1179 ; 3.111 ; 4.500
REMARK 3 NCS RESTRAINTS STATISTICS
REMARK 3 NUMBER OF NCS GROUPS : NULL
REMARK 3
REMARK 3 TLS DETAILS
REMARK 3 NUMBER OF TLS GROUPS : 1
REMARK 3
REMARK 3 TLS GROUP : 1
REMARK 3 NUMBER OF COMPONENTS GROUP : 2
REMARK 3 COMPONENTS C SSSEQI TO C SSSEQI
REMARK 3 RESIDUE RANGE : A 15 A 451
REMARK 3 RESIDUE RANGE : S 795 S 822
REMARK 3 ORIGIN FOR THE GROUP (A): 22.2240 27.6230 28.5830
REMARK 3 T TENSOR
REMARK 3 T11: 0.1744 T22: 0.0216
REMARK 3 T33: 0.0949 T12: -0.0059
REMARK 3 T13: -0.0546 T23: 0.0427
REMARK 3 L TENSOR
REMARK 3 L11: 1.1183 L22: 2.4664
REMARK 3 L33: 1.3415 L12: 0.7934
REMARK 3 L13: 0.5409 L23: 1.2249
REMARK 3 S TENSOR
REMARK 3 S11: 0.0358 S12: -0.1772 S13: -0.0521
REMARK 3 S21: 0.1763 S22: 0.0025 S23: 0.1089
REMARK 3 S31: 0.2114 S32: -0.0339 S33: -0.0383
REMARK 3
REMARK 3 BULK SOLVENT MODELLING.
REMARK 3 METHOD USED : BABINET MODEL WITH MASK
REMARK 3 PARAMETERS FOR MASK CALCULATION
REMARK 3 VDW PROBE RADIUS : 1.40
REMARK 3 ION PROBE RADIUS : 0.80
REMARK 3 SHRINKAGE RADIUS : 0.80
REMARK 3
REMARK 3 OTHER REFINEMENT REMARKS: HYDROGENS HAVE BEEN ADDED IN THE
REMARK 3 RIDING POSITIONS
REMARK 4
REMARK 4 1H2L COMPLIES WITH FORMAT V. 2.3, 09-JULY-1998
REMARK 100
REMARK 100 THIS ENTRY HAS BEEN PROCESSED BY EBI ON 12-AUG-2002.
REMARK 100 THE EBI ID CODE IS EBI-11172.
REMARK 200
REMARK 200 EXPERIMENTAL DETAILS
REMARK 200 EXPERIMENT TYPE : X-RAY DIFFRACTION
REMARK 200 DATE OF DATA COLLECTION : 15-MAY-2002
REMARK 200 TEMPERATURE (KELVIN) : 100
REMARK 200 PH : 7.5
REMARK 200 NUMBER OF CRYSTALS USED : 1
REMARK 200
REMARK 200 SYNCHROTRON (Y/N) : Y
REMARK 200 RADIATION SOURCE : SRS BEAMLINE PX14.2
REMARK 200 BEAMLINE : PX14.2
REMARK 200 X-RAY GENERATOR MODEL : NULL
REMARK 200 MONOCHROMATIC OR LAUE (M/L) : M
REMARK 200 WAVELENGTH OR RANGE (A) : 0.983
REMARK 200 MONOCHROMATOR : NULL
REMARK 200 OPTICS : NULL

REMARK 200
 REMARK 200 DETECTOR TYPE : CCD
 REMARK 200 DETECTOR MANUFACTURER : ADSC
 REMARK 200 INTENSITY-INTEGRATION SOFTWARE : MOSFLM
 REMARK 200 DATA SCALING SOFTWARE : SCALA
 REMARK 200
 REMARK 200 NUMBER OF UNIQUE REFLECTIONS : 27294
 REMARK 200 RESOLUTION RANGE HIGH (A) : 2.25
 REMARK 200 RESOLUTION RANGE LOW (A) : 38.63
 REMARK 200 REJECTION CRITERIA (SIGMA(I)) : NONE
 REMARK 200
 REMARK 200 OVERALL.
 REMARK 200 COMPLETENESS FOR RANGE (%) : 100.0
 REMARK 200 DATA REDUNDANCY : 7.0
 REMARK 200 R MERGE (I) : 0.058
 REMARK 200 R SYM (I) : NULL
 REMARK 200 <I/SIGMA(I)> FOR THE DATA SET : 9.7
 REMARK 200
 REMARK 200 IN THE HIGHEST RESOLUTION SHELL.
 REMARK 200 HIGHEST RESOLUTION SHELL, RANGE HIGH (A) : 2.25
 REMARK 200 HIGHEST RESOLUTION SHELL, RANGE LOW (A) : 2.37
 REMARK 200 COMPLETENESS FOR SHELL (%) : 100.0
 REMARK 200 DATA REDUNDANCY IN SHELL : 7.2
 REMARK 200 R MERGE FOR SHELL (I) : 0.307
 REMARK 200 R SYM FOR SHELL (I) : NULL
 REMARK 200 <I/SIGMA(I)> FOR SHELL : 2.5
 REMARK 200
 REMARK 200 DIFFRACTION PROTOCOL: SINGLE WAVELENGTH
 REMARK 200 METHOD USED TO DETERMINE THE STRUCTURE: MOLECULAR REPLACEMENT
 REMARK 200 SOFTWARE USED: NULL
 REMARK 200 STARTING MODEL: NULL
 REMARK 200
 REMARK 200 REMARK: NULL
 REMARK 280
 REMARK 280 CRYSTAL
 REMARK 280 SOLVENT CONTENT, VS (%): 63
 REMARK 280 MATTHEWS COEFFICIENT, VM (ANGSTROMS**3/DA): 3.4
 REMARK 280
 REMARK 280 CRYSTALLIZATION CONDITIONS: 1.2M AMMONIUM SULPHATE, 4% PEG400,
 REMARK 280 0.1M HEPES PH7.5, ARGON ATMOSPHERE, 11MG/ML PROTEIN WITH
 REMARK 280 1MM FE(II), 2.5MM AKG AND 2.5MM PEPTIDE
 REMARK 290
 REMARK 290 CRYSTALLOGRAPHIC SYMMETRY
 REMARK 290 SYMMETRY OPERATORS FOR SPACE GROUP: P 41 21 2
 REMARK 290

SYMOP	SYMMETRY OPERATOR
1555	X, Y, Z
2555	-X, -Y, 1/2+Z
3555	1/2-Y, 1/2+X, 1/4+Z
4555	1/2+Y, 1/2-X, 3/4+Z
5555	1/2-X, 1/2+Y, 1/4-Z
6555	1/2+X, 1/2-Y, 3/4-Z
7555	Y, X, -Z
8555	-Y, -X, 1/2-Z

 REMARK 290
 REMARK 290 WHERE NNN -> OPERATOR NUMBER
 REMARK 290 MMM -> TRANSLATION VECTOR
 REMARK 290
 REMARK 290 CRYSTALLOGRAPHIC SYMMETRY TRANSFORMATIONS
 REMARK 290 THE FOLLOWING TRANSFORMATIONS OPERATE ON THE ATOM/HETATM

REMARK 290 RECORDS IN THIS ENTRY TO PRODUCE CRYSTALLOGRAPHICALLY
REMARK 290 RELATED MOLECULES.

REMARK 290	SMTRY1	1	1.000000	0.000000	0.000000	0.000000
REMARK 290	SMTRY2	1	0.000000	1.000000	0.000000	0.000000
REMARK 290	SMTRY3	1	0.000000	0.000000	1.000000	0.000000
REMARK 290	SMTRY1	2	-1.000000	0.000000	0.000000	0.000000
REMARK 290	SMTRY2	2	0.000000	-1.000000	0.000000	0.000000
REMARK 290	SMTRY3	2	0.000000	0.000000	1.000000	73.95700
REMARK 290	SMTRY1	3	0.000000	-1.000000	0.000000	43.13200
REMARK 290	SMTRY2	3	1.000000	0.000000	0.000000	43.13200
REMARK 290	SMTRY3	3	0.000000	0.000000	1.000000	36.97850
REMARK 290	SMTRY1	4	0.000000	1.000000	0.000000	43.13200
REMARK 290	SMTRY2	4	-1.000000	0.000000	0.000000	43.13200
REMARK 290	SMTRY3	4	0.000000	0.000000	1.000000	110.93550
REMARK 290	SMTRY1	5	-1.000000	0.000000	0.000000	43.13200
REMARK 290	SMTRY2	5	0.000000	1.000000	0.000000	43.13200
REMARK 290	SMTRY3	5	0.000000	0.000000	-1.000000	36.97850
REMARK 290	SMTRY1	6	1.000000	0.000000	0.000000	43.13200
REMARK 290	SMTRY2	6	0.000000	-1.000000	0.000000	43.13200
REMARK 290	SMTRY3	6	0.000000	0.000000	-1.000000	110.93550
REMARK 290	SMTRY1	7	0.000000	1.000000	0.000000	0.000000
REMARK 290	SMTRY2	7	1.000000	0.000000	0.000000	0.000000
REMARK 290	SMTRY3	7	0.000000	0.000000	-1.000000	0.000000
REMARK 290	SMTRY1	8	0.000000	-1.000000	0.000000	0.000000
REMARK 290	SMTRY2	8	-1.000000	0.000000	0.000000	0.000000
REMARK 290	SMTRY3	8	0.000000	0.000000	-1.000000	73.95700

REMARK 290
REMARK 290 REMARK: NULL
REMARK 300
REMARK 300 BIOMOLECULE: 1
REMARK 300 THIS ENTRY CONTAINS THE CRYSTALLOGRAPHIC ASYMMETRIC UNIT
REMARK 300 WHICH CONSISTS OF 2 CHAIN(S). SEE REMARK 350 FOR
REMARK 300 INFORMATION ON GENERATING THE BIOLOGICAL MOLECULE(S).
REMARK 300
REMARK 300 QUATERNARY STRUCTURE FOR THIS ENTRY: TETRAMERIC
REMARK 300
REMARK 300 THE PROTEIN IS A HOMODIMER FORMED BY CHAIN A.
REMARK 300 A HETERODIMERIC ASSOCIATION OF CHAIN A WITH CHAIN S
REMARK 300 PRODUCES A TETRAMER.
REMARK 300
REMARK 300 THE BURIED SURFACE AREA SHOWN BELOW IS AN AVERAGE
REMARK 300 CALCULATED FOR THE HETEROTETRAMER AND DOES NOT
REMARK 300 CORRESPOND TO THE BURIED SURFACE AREA FOR THE
REMARK 300 HOMODIMER OF CHAIN A
REMARK 300
REMARK 300 THE HETERO-ASSEMBLY DESCRIBED BY REMARK 350 APPEARS
REMARK 300 TO BE A CASE OF STRONG CRYSTAL PACKING WITH
REMARK 300 THE MEAN DIFFERENCE IN ACCESSIBLE SURFACE AREA PER
REMARK 300 CHAIN BETWEEN THE ISOLATED CHAIN AND THAT FOR
REMARK 300 THE CHAIN IN THE COMPLEX IS 2141.3 ANGSTROM**2
REMARK 350
REMARK 350 GENERATING THE BIOMOLECULE
REMARK 350 COORDINATES FOR A COMPLETE MULTIMER REPRESENTING THE KNOWN
REMARK 350 BIOLOGICALLY SIGNIFICANT OLIGOMERIZATION STATE OF THE
REMARK 350 MOLECULE CAN BE GENERATED BY APPLYING BIOMT TRANSFORMATIONS
REMARK 350 GIVEN BELOW. BOTH NON-CRYSTALLOGRAPHIC AND
REMARK 350 CRYSTALLOGRAPHIC OPERATIONS ARE GIVEN.
REMARK 350
REMARK 350 BIOMOLECULE: 1
REMARK 350 APPLY THE FOLLOWING TO CHAINS: A, S
REMARK 350 BIOMT1 1 1.000000 0.000000 0.000000 0.000000

REMARK 350	BIOMT2	1	0.000000	1.000000	0.000000	0.000000
REMARK 350	BIOMT3	1	0.000000	0.000000	1.000000	0.000000
REMARK 350	BIOMT1	2	0.000000	-1.000000	0.000000	86.26400
REMARK 350	BIOMT2	2	-1.000000	0.000000	0.000000	86.26400
REMARK 350	BIOMT3	2	0.000000	0.000000	-1.000000	73.95700

REMARK 465 MISSING RESIDUES

REMARK 465 THE FOLLOWING RESIDUES WERE NOT LOCATED IN THE
 REMARK 465 EXPERIMENT. (M=MODEL NUMBER; RES=RESIDUE NAME; C=CHAIN
 REMARK 465 IDENTIFIER; SSSEQ=SEQUENCE NUMBER; I=INSERTION CODE.)

REMARK 465	M	RES	C	SSSEQI
REMARK 465		MET	A	1
REMARK 465		ALA	A	2
REMARK 465		ALA	A	3
REMARK 465		THR	A	4
REMARK 465		ALA	A	5
REMARK 465		ALA	A	6
REMARK 465		GLU	A	7
REMARK 465		ALA	A	8
REMARK 465		VAL	A	9
REMARK 465		ALA	A	10
REMARK 465		SER	A	11
REMARK 465		GLY	A	12
REMARK 465		SER	A	13
REMARK 465		GLY	A	14
REMARK 465		LYS	A	304
REMARK 465		ARG	A	305
REMARK 465		ILE	A	306
REMARK 465		SER	S	786
REMARK 465		MET	S	787
REMARK 465		ASP	S	788
REMARK 465		GLU	S	789
REMARK 465		SER	S	790
REMARK 465		GLY	S	791
REMARK 465		LEU	S	792
REMARK 465		PRO	S	793
REMARK 465		GLN	S	794
REMARK 465		GLN	S	807
REMARK 465		GLY	S	808
REMARK 465		SER	S	809
REMARK 465		ARG	S	810
REMARK 465		ASN	S	811
REMARK 465		LEU	S	812
REMARK 465		ASP	S	823
REMARK 465		GLN	S	824
REMARK 465		VAL	S	825
REMARK 465		ASN	S	826

REMARK 470

REMARK 470 MISSING ATOM

REMARK 470 THE FOLLOWING RESIDUES HAVE MISSING ATOMS (M=MODEL NUMBER;
 REMARK 470 RES=RESIDUE NAME; C=CHAIN IDENTIFIER; SSEQ=SEQUENCE NUMBER;
 REMARK 470 I=INSERTION CODE):

REMARK 470	M	RES	C	SSEQI	ATOMS
REMARK 470		GLU	A	15	CG CD OE1 OE2
REMARK 470		GLU	A	29	CG CD OE1 OE2
REMARK 470		ASN	A	87	CG OD1 ND2
REMARK 470		LYS	A	106	CD CE NZ
REMARK 470		ARG	A	117	CG CD NE CZ NH1 NH2
REMARK 470		GLN	A	133	CG CD OE1 NE2
REMARK 470		GLN	A	136	CG CD OE1 NE2

REMARK 470 GLN A 137 CG CD OE1 NE2
 REMARK 470 ARG A 156 CG CD NE CZ NH1 NH2
 REMARK 470 LYS A 157 CD CE NZ
 REMARK 470 LYS A 311 CG CD CE NZ
 REMARK 500
 REMARK 500 GEOMETRY AND STEREOCHEMISTRY
 REMARK 500 SUBTOPIC: COVALENT BOND LENGTHS
 REMARK 500
 REMARK 500 THE STEREOCHEMICAL PARAMETERS OF THE FOLLOWING RESIDUES
 REMARK 500 HAVE VALUES WHICH DEVIATE FROM EXPECTED VALUES BY MORE
 REMARK 500 THAN 6*RMSD AND BY MORE THAN 0.150 ANGSTROMS (M=MODEL
 REMARK 500 NUMBER; RES=RESIDUE NAME; C=CHAIN IDENTIFIER; SSEQ=SEQUENCE
 REMARK 500 NUMBER; I=INSERTION CODE).
 REMARK 500
 REMARK 500 STANDARD TABLE:
 REMARK 500 FORMAT: (10X,I3,1X,A3,1X,A1,I4,A1,1X,2(A4,A1,3X),12X,F5.3)
 REMARK 500
 REMARK 500 EXPECTED VALUESS: ENGH AND HUBER, 1991
 REMARK 500
 REMARK 500 M RES CSSEQI ATM1 RES CSSEQI ATM2 DEVIATION
 REMARK 500 MET A 343 SD MET A 343 CE -0.243
 REMARK 500
 REMARK 500 REMARK: NULL
 REMARK 500
 REMARK 500 GEOMETRY AND STEREOCHEMISTRY
 REMARK 500 SUBTOPIC: CLOSE CONTACTS IN SAME ASYMMETRIC UNIT
 REMARK 500
 REMARK 500 THE FOLLOWING ATOMS ARE IN CLOSE CONTACT.
 REMARK 500

ATM1	RES	C	SSEQI	ATM2	RES	C	SSEQI	DISTANCE
O	ALA	A	300	OH	TYR	S	798	2.18

 REMARK 500
 REMARK 525
 REMARK 525 SOLVENT
 REMARK 525
 REMARK 525 THE SOLVENT MOLECULES ARE GIVEN CHAIN IDENTIFIERS TO
 REMARK 525 INDICATE THE PROTEIN CHAIN TO WHICH THEY ARE MOST CLOSELY
 REMARK 525 ASSOCIATED WITH:

PROTEIN CHAIN	SOLVENT CHAIN
A	Z
S	H

 REMARK 525
 REMARK 600
 REMARK 600 HETEROGEN
 REMARK 600
 REMARK 600 FOR METAL ATOM FE FE2 A1350 THE COORDINATION ANGLES ARE:

	1	2	3	4
1 HIS 199A NE2				
2 ASP 201A OD2	103.4			
3 HIS 279A NE2	83.1	86.0		
4 AKG 1351A O1	168.9	87.6	98.2	
5 AKG 1351A O5	87.0	169.4	97.2	81.9

 REMARK 600
 REMARK 700
 REMARK 700 SHEET
 REMARK 700 THE SHEET STRUCTURE OF THIS MOLECULE IS BIFURCATED. IN
 REMARK 700 ORDER TO REPRESENT THIS FEATURE IN THE SHEET RECORDS BELOW,
 REMARK 700 TWO SHEETS ARE DEFINED.
 REMARK 800
 REMARK 800 SITE
 REMARK 800 SITE_IDENTIFIER: FE1
 REMARK 800 SITE_DESCRIPTION: FE BINDING SITE FOR CHAIN A
 REMARK 800

REMARK 800 SITE_IDENTIFIER: AKG
 REMARK 800 SITE_DESCRIPTION: AKG BINDING SITE FOR CHAIN A
 REMARK 800
 REMARK 800 SITE_IDENTIFIER: SO1
 REMARK 800 SITE_DESCRIPTION: SO4 BINDING SITE FOR CHAIN A
 REMARK 800
 REMARK 800 SITE_IDENTIFIER: SO2
 REMARK 800 SITE_DESCRIPTION: SO4 BINDING SITE FOR CHAIN A
 REMARK 900
 REMARK 900 RELATED ENTRIES
 REMARK 900 RELATED ID: 1D7G RELATED DB: PDB
 REMARK 900 A MODEL FOR THE COMPLEX BETWEEN THE
 REMARK 900 HYPOXIA-INDUCIBLE FACTOR-1 (HIF-1) AND ITS
 REMARK 900 CONSENSUS DEOXYRIBONUCLEIC ACID SEQUENCE
 REMARK 900 RELATED ID: 1H2K RELATED DB: PDB
 REMARK 900 FACTOR INHIBITING HIF-1 ALPHA IN COMPLEX
 REMARK 900 WITH HIF-1 ALPHA FRAGMENT PEPTIDE
 REMARK 900 RELATED ID: 1H2M RELATED DB: PDB
 REMARK 900 FACTOR INHIBITING HIF-1 ALPHA IN COMPLEX
 REMARK 900 WITH HIF-1 ALPHA FRAGMENT PEPTIDE
 REMARK 900 RELATED ID: 1H2N RELATED DB: PDB
 REMARK 900 FACTOR INHIBITING HIF-1 ALPHA IN COMPLEX
 REMARK 900 WITH HIF-1 ALPHA FRAGMENT PEPTIDE
 REMARK 900 RELATED ID: 1L8C RELATED DB: PDB
 REMARK 900 STRUCTURAL BASIS FOR HIF-1ALPHA/CBP
 REMARK 900 RECOGNITION IN THECELLULAR HYPOXIC RESPONSE
 REMARK 900 RELATED ID: 1LM8 RELATED DB: PDB
 REMARK 900 STRUCTURE OF A HIF-1A-PVHL-ELONGINB-
 REMARK 900 ELONGINC COMPLEX
 REMARK 900 RELATED ID: 1LQB RELATED DB: PDB
 REMARK 900 CRYSTAL STRUCTURE OF A HYDROXYLATED HIF-1
 REMARK 900 ALPHA PEPTIDEBOUND TO THE PVHL/ELONGIN-C/
 REMARK 900 ELONGIN-B COMPLEX
 DBREF 1H2L A 1 349 SWS Q969Q7 Q969Q7 1 349
 DBREF 1H2L S 786 826 SWS Q16665 H1FA_HUMAN 786 826
 SEQRES 1 A 349 MET ALA ALA THR ALA ALA GLU ALA VAL ALA SER GLY SER
 SEQRES 2 A 349 GLY GLU PRO ARG GLU GLU ALA GLY ALA LEU GLY PRO ALA
 SEQRES 3 A 349 TRP ASP GLU SER GLN LEU ARG SER TYR SER PHE PRO THR
 SEQRES 4 A 349 ARG PRO ILE PRO ARG LEU SER GLN SER ASP PRO ARG ALA
 SEQRES 5 A 349 GLU GLU LEU ILE GLU ASN GLU GLU PRO VAL VAL LEU THR
 SEQRES 6 A 349 ASP THR ASN LEU VAL TYR PRO ALA LEU LYS TRP ASP LEU
 SEQRES 7 A 349 GLU TYR LEU GLN GLU ASN ILE GLY ASN GLY ASP PHE SER
 SEQRES 8 A 349 VAL TYR SER ALA SER THR HIS LYS PHE LEU TYR TYR ASP
 SEQRES 9 A 349 GLU LYS LYS MET ALA ASN PHE GLN ASN PHE LYS PRO ARG
 SEQRES 10 A 349 SER ASN ARG GLU GLU MET LYS PHE HIS GLU PHE VAL GLU
 SEQRES 11 A 349 LYS LEU GLN ASP ILE GLN GLN ARG GLY GLY GLU GLU ARG
 SEQRES 12 A 349 LEU TYR LEU GLN GLN THR LEU ASN ASP THR VAL GLY ARG
 SEQRES 13 A 349 LYS ILE VAL MET ASP PHE LEU GLY PHE ASN TRP ASN TRP
 SEQRES 14 A 349 ILE ASN LYS GLN GLN GLY LYS ARG GLY TRP GLY GLN LEU
 SEQRES 15 A 349 THR SER ASN LEU LEU LEU ILE GLY MET GLU GLY ASN VAL
 SEQRES 16 A 349 THR PRO ALA HIS TYR ASP GLU GLN GLN ASN PHE PHE ALA
 SEQRES 17 A 349 GLN ILE LYS GLY TYR LYS ARG CYS ILE LEU PHE PRO PRO
 SEQRES 18 A 349 ASP GLN PHE GLU CYS LEU TYR PRO TYR PRO VAL HIS HIS
 SEQRES 19 A 349 PRO CYS ASP ARG GLN SER GLN VAL ASP PHE ASP ASN PRO
 SEQRES 20 A 349 ASP TYR GLU ARG PHE PRO ASN PHE GLN ASN VAL VAL GLY
 SEQRES 21 A 349 TYR GLU THR VAL VAL GLY PRO GLY ASP VAL LEU TYR ILE
 SEQRES 22 A 349 PRO MET TYR TRP TRP HIS HIS ILE GLU SER LEU LEU ASN
 SEQRES 23 A 349 GLY GLY ILE THR ILE THR VAL ASN PHE TRP TYR LYS GLY
 SEQRES 24 A 349 ALA PRO THR PRO LYS ARG ILE GLU TYR PRO LEU LYS ALA
 SEQRES 25 A 349 HIS GLN LYS VAL ALA ILE MET ARG ASN ILE GLU LYS MET
 SEQRES 26 A 349 LEU GLY GLU ALA LEU GLY ASN PRO GLN GLU VAL GLY PRO

```

SEQRES 27 A 349 LEU LEU ASN THR MET ILE LYS GLY ARG TYR ASN
SEQRES 1 S 41 SER MET ASP GLU SER GLY LEU PRO GLN LEU THR SER TYR
SEQRES 2 S 41 ASP CYS GLU VAL ASN ALA PRO ILE GLN GLY SER ARG ASN
SEQRES 3 S 41 LEU LEU GLN GLY GLU GLU LEU LEU ARG ALA LEU ASP GLN
SEQRES 4 S 41 VAL ASN
HET FE2 A1350 1
HET AKG A1351 10
HET SO4 A1352 5
HET SO4 A1353 5
HETNAM FE2 FE (II) ION
HETNAM AKG 2-OXYGLUTARIC ACID
HETNAM SO4 SULFATE ION
FORMUL 3 FE2 FE1 2+
FORMUL 4 AKG C5 H6 O5
FORMUL 5 SO4 2(O4 S1 2-)
FORMUL 6 HOH *139(H2 O1)
HELIX 1 1 ASP A 28 LEU A 32 5
HELIX 2 2 ASP A 49 ASN A 58 1
HELIX 3 3 VAL A 70 TRP A 76 5
HELIX 4 4 ASP A 77 ILE A 85 1
HELIX 5 5 ASP A 104 PHE A 111 5
HELIX 6 6 LYS A 124 ARG A 138 1
HELIX 7 7 GLY A 155 GLY A 164 1
HELIX 8 8 ASN A 166 GLY A 178 1
HELIX 9 9 PRO A 220 ASP A 222 5
HELIX 10 10 GLN A 223 TYR A 228 1
HELIX 11 11 PHE A 252 VAL A 258 5
HELIX 12 12 LYS A 311 GLY A 331 1
HELIX 13 13 ASN A 332 GLN A 334 5
HELIX 14 14 GLU A 335 LYS A 345 1
HELIX 15 15 GLN S 814 LEU S 822 1
SHEET 1 AA 5 THR A 39 PRO A 41 0
SHEET 2 AA 5 GLY A 260 VAL A 265 1 O GLY A 260 N ARG A 40
SHEET 3 AA 5 LYS A 214 PHE A 219 -1 O LYS A 214 N VAL A 265
SHEET 4 AA 5 TRP A 278 SER A 283 -1 O TRP A 278 N PHE A 219
SHEET 5 AA 5 VAL A 195 HIS A 199 -1 O THR A 196 N ILE A 281
SHEET 1 AB 6 ARG A 44 LEU A 45 0
SHEET 2 AB 6 VAL A 62 LEU A 64 1 O VAL A 63 N LEU A 45
SHEET 3 AB 6 VAL A 270 ILE A 273 -1 O VAL A 270 N LEU A 64
SHEET 4 AB 6 GLN A 204 LYS A 211 -1 O ASN A 205 N ILE A 273
SHEET 5 AB 6 THR A 290 TYR A 297 -1 O ILE A 291 N ILE A 210
SHEET 6 AB 6 LEU A 182 SER A 184 -1 N THR A 183 O TRP A 296
SHEET 1 AC 9 ARG A 44 LEU A 45 0
SHEET 2 AC 9 VAL A 62 LEU A 64 1 O VAL A 63 N LEU A 45
SHEET 3 AC 9 VAL A 270 ILE A 273 -1 O VAL A 270 N LEU A 64
SHEET 4 AC 9 GLN A 204 LYS A 211 -1 O ASN A 205 N ILE A 273
SHEET 5 AC 9 THR A 290 TYR A 297 -1 O ILE A 291 N ILE A 210
SHEET 6 AC 9 LEU A 186 GLY A 190 -1 O LEU A 186 N ASN A 294
SHEET 7 AC 9 ARG A 143 THR A 149 -1 O LEU A 146 N ILE A 189
SHEET 8 AC 9 PHE A 90 ALA A 95 -1 O SER A 91 N GLN A 147
SHEET 9 AC 9 SER A 118 MET A 123 -1 O ASN A 119 N SER A 94
LINK FE FE2 A1350 NE2 HIS A 199 1555 1555
LINK FE FE2 A1350 OD2 ASP A 201 1555 1555
LINK FE FE2 A1350 NE2 HIS A 279 1555 1555
LINK FE FE2 A1350 O1 AKG A1351 1555 1555
LINK FE FE2 A1350 O5 AKG A1351 1555 1555
CISPEP 1 TYR A 308 PRO A 309 0 1.27
SITE 1 FE1 3 HIS A 199 ASP A 201 HIS A 279
SITE 1 AKG 12 TYR A 145 THR A 196 HIS A 199 ASP A 201
SITE 2 AKG 12 ASN A 205 PHE A 207 LYS A 214 HIS A 279
SITE 3 AKG 12 ILE A 281 ASN A 294 TRP A 296 HOH Z 65

```

SITE	1	SO1	4	ARG	A	138	GLY	A	140	GLU	A	141	GLU	A	142	
SITE	1	SO2	5	ARG	A	143	GLU	A	192	GLY	A	193	LEU	A	285	
SITE	2	SO2	5	ASN	A	286										
CRYST1	86.264		86.264		147.914		90.00		90.00		90.00		P	41	21	2
ORIGX1	1.000000	0.000000	0.000000	0.000000						0.000000						8
ORIGX2	0.000000	1.000000	0.000000	0.000000						0.000000						
ORIGX3	0.000000	0.000000	1.000000	0.000000						0.000000						
SCALE1	0.011592	0.000000	0.000000	0.000000						0.000000						
SCALE2	0.000000	0.011592	0.000000	0.000000						0.000000						
SCALE3	0.000000	0.000000	0.006761	0.000000						0.000000						
ATOM	1	N		GLU	A	15	8.505	32.866		9.893	1.00	61.72				
ATOM	2	CA		GLU	A	15	7.173	32.223		9.682	1.00	61.95				
ATOM	3	C		GLU	A	15	7.251	30.749	10.070		1.00	61.54				
ATOM	4	O		GLU	A	15	8.109	30.026	9.581	1.00	61.66					
ATOM	5	CB		GLU	A	15	6.724	32.375	8.234	1.00	62.08					
ATOM	6	N		PRO	A	16	6.353	30.305	10.941	1.00	61.24					
ATOM	7	CA		PRO	A	16	6.386	28.928	11.455	1.00	60.97					
ATOM	8	C		PRO	A	16	6.342	27.853	10.368	1.00	60.27					
ATOM	9	O		PRO	A	16	5.494	27.879	9.479	1.00	60.01					
ATOM	10	CB		PRO	A	16	5.134	28.849	12.333	1.00	61.10					
ATOM	11	CG		PRO	A	16	4.790	30.262	12.653	1.00	61.53					
ATOM	12	CD		PRO	A	16	5.228	31.079	11.488	1.00	61.35					
ATOM	13	N		ARG	A	17	7.256	26.900	10.464	1.00	59.45					
ATOM	14	CA		ARG	A	17	7.348	25.820	9.494	1.00	59.05					
ATOM	15	C		ARG	A	17	6.083	24.971	9.478	1.00	57.86					
ATOM	16	O		ARG	A	17	5.312	24.963	10.437	1.00	57.86					
ATOM	17	CB		ARG	A	17	8.541	24.913	9.827	1.00	59.53					
ATOM	18	CG		ARG	A	17	9.906	25.621	9.859	1.00	60.69					
ATOM	19	CD		ARG	A	17	11.080	24.696	10.194	1.00	62.32					
ATOM	20	NE		ARG	A	17	11.040	24.185	11.567	1.00	63.73					
ATOM	21	CZ		ARG	A	17	11.464	24.852	12.649	1.00	65.28					
ATOM	22	NH1		ARG	A	17	11.962	26.085	12.544	1.00	65.01					
ATOM	23	NH2		ARG	A	17	11.385	24.282	13.848	1.00	65.09					
ATOM	24	N		GLU	A	18	5.878	24.254	8.382	1.00	56.26					
ATOM	25	CA		GLU	A	18	4.749	23.351	8.273	1.00	55.12					
ATOM	26	C		GLU	A	18	5.222	21.931	8.587	1.00	53.54					
ATOM	27	O		GLU	A	18	6.214	21.468	8.031	1.00	52.86					
ATOM	28	CB		GLU	A	18	4.150	23.406	6.867	1.00	55.33					
ATOM	29	CG		GLU	A	18	3.482	24.731	6.519	1.00	56.39					
ATOM	30	CD		GLU	A	18	2.100	24.904	7.137	1.00	57.19					
ATOM	31	OE1		GLU	A	18	1.559	23.935	7.718	1.00	56.35					
ATOM	32	OE2		GLU	A	18	1.548	26.023	7.028	1.00	58.12					
ATOM	33	N		GLU	A	19	4.526	21.260	9.501	1.00	51.73					
ATOM	34	CA		GLU	A	19	4.823	19.867	9.816	1.00	50.56					
ATOM	35	C		GLU	A	19	4.409	18.944	8.663	1.00	48.36					
ATOM	36	O		GLU	A	19	3.312	19.066	8.102	1.00	47.54					
ATOM	37	CB		GLU	A	19	4.115	19.422	11.100	1.00	51.05					
ATOM	38	CG		GLU	A	19	4.577	20.139	12.360	1.00	53.53					
ATOM	39	CD		GLU	A	19	4.363	19.323	13.638	1.00	57.67					
ATOM	40	OE1		GLU	A	19	3.906	18.146	13.561	1.00	59.24					
ATOM	41	OE2		GLU	A	19	4.663	19.864	14.735	1.00	59.50					
ATOM	42	N		ALA	A	20	5.314	18.035	8.320	1.00	46.04					
ATOM	43	CA		ALA	A	20	5.100	17.030	7.283	1.00	44.66					
ATOM	44	C		ALA	A	20	3.741	16.365	7.394	1.00	42.92					
ATOM	45	O		ALA	A	20	3.230	16.154	8.491	1.00	42.14					
ATOM	46	CB		ALA	A	20	6.182	15.965	7.360	1.00	44.46					
ATOM	47	N		GLY	A	21	3.173	16.025	6.247	1.00	41.16					
ATOM	48	CA		GLY	A	21	1.897	15.341	6.215	1.00	40.22					
ATOM	49	C		GLY	A	21	0.757	16.300	6.480	1.00	39.27					
ATOM	50	O		GLY	A	21	-0.309	15.895	6.908	1.00	38.03					
ATOM	51	N		ALA	A	22	1.005	17.584	6.240	1.00	38.86					

N
C
C
O
C
N
C
C
O
C
C
N
C
C
C
O
C
C
N
N
N
N
C
C
O
C
C
C
O
O
N
C
C
O
C
C
O
O
N
C
C
O
C
N
C
C
O
N

ATOM	52	CA	ALA	A	22	-0.018	18.611	6.371	1.00	38.79	C
ATOM	53	C	ALA	A	22	-0.618	18.613	7.758	1.00	38.64	C
ATOM	54	O	ALA	A	22	-1.820	18.759	7.924	1.00	38.07	O
ATOM	55	CB	ALA	A	22	-1.102	18.409	5.324	1.00	38.68	C
ATOM	56	N	LEU	A	23	0.231	18.451	8.760	1.00	39.07	N
ATOM	57	CA	LEU	A	23	-0.233	18.473	10.135	1.00	39.72	C
ATOM	58	C	LEU	A	23	-0.290	19.886	10.692	1.00	39.39	C
ATOM	59	O	LEU	A	23	-0.464	20.075	11.880	1.00	39.41	O
ATOM	60	CB	LEU	A	23	0.642	17.574	11.003	1.00	40.11	C
ATOM	61	CG	LEU	A	23	0.528	16.118	10.558	1.00	41.82	C
ATOM	62	CD1	LEU	A	23	1.414	15.185	11.384	1.00	43.18	C
ATOM	63	CD2	LEU	A	23	-0.916	15.673	10.633	1.00	43.43	C
ATOM	64	N	GLY	A	24	-0.156	20.879	9.823	1.00	39.21	N
ATOM	65	CA	GLY	A	24	-0.290	22.258	10.237	1.00	39.16	C
ATOM	66	C	GLY	A	24	0.964	22.862	10.820	1.00	39.21	C
ATOM	67	O	GLY	A	24	2.011	22.219	10.916	1.00	39.16	O
ATOM	68	N	PRO	A	25	0.854	24.118	11.223	1.00	39.08	N
ATOM	69	CA	PRO	A	25	1.997	24.830	11.775	1.00	38.84	C
ATOM	70	C	PRO	A	25	2.410	24.197	13.086	1.00	38.68	C
ATOM	71	O	PRO	A	25	1.572	23.842	13.914	1.00	38.08	O
ATOM	72	CB	PRO	A	25	1.477	26.259	12.001	1.00	38.76	C
ATOM	73	CG	PRO	A	25	-0.004	26.197	11.895	1.00	39.12	C
ATOM	74	CD	PRO	A	25	-0.375	24.930	11.210	1.00	39.17	C
ATOM	75	N	ALA	A	26	3.713	24.037	13.246	1.00	38.92	N
ATOM	76	CA	ALA	A	26	4.269	23.489	14.467	1.00	39.38	C
ATOM	77	C	ALA	A	26	3.924	24.381	15.676	1.00	38.73	C
ATOM	78	O	ALA	A	26	3.677	23.882	16.772	1.00	39.39	O
ATOM	79	CB	ALA	A	26	5.762	23.347	14.319	1.00	39.60	C
ATOM	80	N	TRP	A	27	3.871	25.687	15.468	1.00	37.43	N
ATOM	81	CA	TRP	A	27	3.516	26.602	16.543	1.00	36.82	C
ATOM	82	C	TRP	A	27	3.034	27.919	15.941	1.00	36.10	C
ATOM	83	O	TRP	A	27	3.013	28.074	14.731	1.00	35.54	O
ATOM	84	CB	TRP	A	27	4.746	26.835	17.424	1.00	36.91	C
ATOM	85	CG	TRP	A	27	5.949	27.019	16.596	1.00	36.01	C
ATOM	86	CD1	TRP	A	27	6.770	26.041	16.089	1.00	36.29	C
ATOM	87	CD2	TRP	A	27	6.450	28.249	16.107	1.00	34.41	C
ATOM	88	NE1	TRP	A	27	7.761	26.610	15.324	1.00	36.06	N
ATOM	89	CE2	TRP	A	27	7.592	27.966	15.332	1.00	34.97	C
ATOM	90	CE3	TRP	A	27	6.066	29.568	16.263	1.00	34.24	C
ATOM	91	CZ2	TRP	A	27	8.332	28.947	14.720	1.00	36.39	C
ATOM	92	CZ3	TRP	A	27	6.808	30.539	15.664	1.00	36.24	C
ATOM	93	CH2	TRP	A	27	7.927	30.225	14.890	1.00	36.25	C
ATOM	94	N	ASP	A	28	2.620	28.860	16.775	1.00	35.59	N
ATOM	95	CA	ASP	A	28	2.253	30.168	16.267	1.00	35.15	C
ATOM	96	C	ASP	A	28	2.816	31.224	17.160	1.00	33.93	C
ATOM	97	O	ASP	A	28	3.311	30.948	18.250	1.00	33.56	O
ATOM	98	CB	ASP	A	28	0.739	30.338	16.131	1.00	36.08	C
ATOM	99	CG	ASP	A	28	0.038	30.310	17.449	1.00	38.11	C
ATOM	100	OD1	ASP	A	28	-0.096	31.392	18.076	1.00	41.06	O
ATOM	101	OD2	ASP	A	28	-0.392	29.240	17.938	1.00	41.04	O
ATOM	102	N	GLU	A	29	2.737	32.448	16.665	1.00	32.84	N
ATOM	103	CA	GLU	A	29	3.288	33.615	17.330	1.00	31.31	C
ATOM	104	C	GLU	A	29	2.792	33.767	18.756	1.00	29.80	C
ATOM	105	O	GLU	A	29	3.547	34.124	19.639	1.00	29.03	O
ATOM	106	CB	GLU	A	29	2.938	34.841	16.537	1.00	31.81	C
ATOM	107	N	SER	A	30	1.524	33.478	18.989	1.00	28.55	N
ATOM	108	CA	SER	A	30	0.957	33.680	20.314	1.00	28.00	C
ATOM	109	C	SER	A	30	1.613	32.830	21.391	1.00	27.25	C
ATOM	110	O	SER	A	30	1.360	33.035	22.563	1.00	26.88	O
ATOM	111	CB	SER	A	30	-0.546	33.404	20.302	1.00	27.71	C
ATOM	112	OG	SER	A	30	-0.800	32.015	20.328	1.00	28.97	O

ATOM	113	N	GLN	A	31	2.434	31.864	20.994	1.00	26.92
ATOM	114	CA	GLN	A	31	3.089	30.988	21.950	1.00	26.85
ATOM	115	C	GLN	A	31	4.447	31.558	22.362	1.00	26.63
ATOM	116	O	GLN	A	31	5.115	30.995	23.220	1.00	25.89
ATOM	117	CB	GLN	A	31	3.270	29.576	21.375	1.00	26.90
ATOM	118	CG	GLN	A	31	1.975	28.816	21.097	1.00	27.50
ATOM	119	CD	GLN	A	31	2.227	27.446	20.480	1.00	27.30
ATOM	120	OE1	GLN	A	31	2.332	27.330	19.262	1.00	28.39
ATOM	121	NE2	GLN	A	31	2.354	26.419	21.319	1.00	23.43
ATOM	122	N	LEU	A	32	4.835	32.682	21.757	1.00	26.56
ATOM	123	CA	LEU	A	32	6.094	33.351	22.078	1.00	26.88
ATOM	124	C	LEU	A	32	5.854	34.500	23.047	1.00	26.64
ATOM	125	O	LEU	A	32	4.875	35.214	22.913	1.00	26.22
ATOM	126	CB	LEU	A	32	6.743	33.906	20.807	1.00	26.71
ATOM	127	CG	LEU	A	32	7.054	32.866	19.724	1.00	29.00
ATOM	128	CD1	LEU	A	32	7.704	33.537	18.535	1.00	31.11
ATOM	129	CD2	LEU	A	32	7.942	31.773	20.245	1.00	28.07
ATOM	130	N	ARG	A	33	6.737	34.671	24.026	1.00	26.49
ATOM	131	CA	ARG	A	33	6.622	35.799	24.957	1.00	26.48
ATOM	132	C	ARG	A	33	7.070	37.071	24.252	1.00	26.68
ATOM	133	O	ARG	A	33	7.810	37.025	23.280	1.00	26.97
ATOM	134	CB	ARG	A	33	7.454	35.554	26.224	1.00	26.44
ATOM	135	CG	ARG	A	33	7.071	34.261	26.976	1.00	25.53
ATOM	136	CD	ARG	A	33	7.869	34.029	28.245	1.00	25.76
ATOM	137	NE	ARG	A	33	7.329	32.926	29.021	1.00	25.87
ATOM	138	CZ	ARG	A	33	6.418	33.033	29.969	1.00	25.35
ATOM	139	NH1	ARG	A	33	5.916	34.198	30.309	1.00	24.02
ATOM	140	NH2	ARG	A	33	6.003	31.941	30.587	1.00	28.18
ATOM	141	N	SER	A	34	6.643	38.214	24.751	1.00	26.73
ATOM	142	CA	SER	A	34	6.939	39.469	24.090	1.00	27.21
ATOM	143	C	SER	A	34	8.021	40.251	24.840	1.00	26.40
ATOM	144	O	SER	A	34	7.957	40.391	26.046	1.00	25.65
ATOM	145	CB	SER	A	34	5.657	40.278	24.028	1.00	27.92
ATOM	146	OG	SER	A	34	5.402	40.780	25.323	1.00	31.78
ATOM	147	N	TYR	A	35	9.009	40.750	24.110	1.00	26.09
ATOM	148	CA	TYR	A	35	10.169	41.390	24.711	1.00	26.44
ATOM	149	C	TYR	A	35	10.412	42.731	24.046	1.00	27.00
ATOM	150	O	TYR	A	35	9.815	43.009	23.028	1.00	27.14
ATOM	151	CB	TYR	A	35	11.386	40.479	24.577	1.00	25.83
ATOM	152	CG	TYR	A	35	11.217	39.198	25.357	1.00	25.49
ATOM	153	CD1	TYR	A	35	11.041	39.226	26.739	1.00	23.65
ATOM	154	CD2	TYR	A	35	11.219	37.963	24.723	1.00	23.87
ATOM	155	CE1	TYR	A	35	10.869	38.063	27.459	1.00	23.41
ATOM	156	CE2	TYR	A	35	11.061	36.795	25.445	1.00	24.37
ATOM	157	CZ	TYR	A	35	10.881	36.847	26.809	1.00	22.92
ATOM	158	OH	TYR	A	35	10.698	35.686	27.522	1.00	23.94
ATOM	159	N	SER	A	36	11.326	43.531	24.596	1.00	27.45
ATOM	160	CA	SER	A	36	11.555	44.905	24.127	1.00	27.58
ATOM	161	C	SER	A	36	12.553	45.092	22.987	1.00	27.13
ATOM	162	O	SER	A	36	12.764	46.211	22.533	1.00	27.71
ATOM	163	CB	SER	A	36	12.109	45.724	25.286	1.00	27.62
ATOM	164	OG	SER	A	36	13.365	45.201	25.697	1.00	27.53
ATOM	165	N	PHE	A	37	13.181	44.025	22.543	1.00	25.62
ATOM	166	CA	PHE	A	37	14.263	44.169	21.590	1.00	24.95
ATOM	167	C	PHE	A	37	13.949	43.447	20.301	1.00	25.32
ATOM	168	O	PHE	A	37	13.191	42.512	20.274	1.00	25.29
ATOM	169	CB	PHE	A	37	15.537	43.571	22.185	1.00	23.88
ATOM	170	CG	PHE	A	37	15.340	42.162	22.731	1.00	22.69
ATOM	171	CD1	PHE	A	37	15.353	41.077	21.890	1.00	20.43
ATOM	172	CD2	PHE	A	37	15.104	41.949	24.076	1.00	23.12
ATOM	173	CE1	PHE	A	37	15.160	39.778	22.386	1.00	22.83

ATOM	174	CE2	PHE	A	37	14.920	40.666	24.578	1.00	22.73	
ATOM	175	CZ	PHE	A	37	14.944	39.585	23.737	1.00	22.04	C
ATOM	176	N	PRO	A	38	14.533	43.907	19.222	1.00	25.57	C
ATOM	177	CA	PRO	A	38	14.396	43.216	17.950	1.00	25.50	C
ATOM	178	C	PRO	A	38	15.321	42.002	17.861	1.00	25.35	C
ATOM	179	O	PRO	A	38	16.280	41.887	18.640	1.00	25.01	O
ATOM	180	CB	PRO	A	38	14.815	44.278	16.944	1.00	25.31	C
ATOM	181	CG	PRO	A	38	15.776	45.181	17.729	1.00	26.61	C
ATOM	182	CD	PRO	A	38	15.271	45.178	19.125	1.00	25.73	C
ATOM	183	N	THR	A	39	15.032	41.124	16.902	1.00	24.60	N
ATOM	184	CA	THR	A	39	15.876	39.979	16.607	1.00	24.58	C
ATOM	185	C	THR	A	39	15.881	39.686	15.112	1.00	25.39	C
ATOM	186	O	THR	A	39	15.004	40.121	14.383	1.00	24.72	O
ATOM	187	CB	THR	A	39	15.364	38.721	17.295	1.00	24.33	C
ATOM	188	OG1	THR	A	39	14.023	38.453	16.860	1.00	21.36	O
ATOM	189	CG2	THR	A	39	15.256	38.892	18.832	1.00	23.91	C
ATOM	190	N	ARG	A	40	16.875	38.912	14.692	1.00	26.09	N
ATOM	191	CA	ARG	A	40	17.000	38.399	13.337	1.00	26.92	C
ATOM	192	C	ARG	A	40	17.057	36.884	13.469	1.00	26.73	C
ATOM	193	O	ARG	A	40	17.407	36.366	14.517	1.00	26.53	O
ATOM	194	CB	ARG	A	40	18.291	38.888	12.696	1.00	27.12	C
ATOM	195	CG	ARG	A	40	18.289	40.360	12.400	1.00	32.37	C
ATOM	196	CD	ARG	A	40	16.925	40.858	11.994	1.00	36.79	C
ATOM	197	NE	ARG	A	40	16.783	41.137	10.583	1.00	41.20	N
ATOM	198	CZ	ARG	A	40	15.617	41.408	10.023	1.00	46.28	C
ATOM	199	NH1	ARG	A	40	14.512	41.388	10.768	1.00	47.83	N
ATOM	200	NH2	ARG	A	40	15.548	41.708	8.731	1.00	48.02	N
ATOM	201	N	PRO	A	41	16.728	36.163	12.413	1.00	26.96	N
ATOM	202	CA	PRO	A	41	16.709	34.706	12.510	1.00	27.17	N
ATOM	203	C	PRO	A	41	18.085	34.054	12.469	1.00	26.50	C
ATOM	204	O	PRO	A	41	19.002	34.538	11.830	1.00	26.34	O
ATOM	205	CB	PRO	A	41	15.867	34.277	11.298	1.00	27.12	C
ATOM	206	CG	PRO	A	41	16.027	35.434	10.296	1.00	27.66	C
ATOM	207	CD	PRO	A	41	16.337	36.661	11.077	1.00	26.70	C
ATOM	208	N	ILE	A	42	18.214	32.951	13.185	1.00	25.35	N
ATOM	209	CA	ILE	A	42	19.400	32.143	13.070	1.00	24.29	C
ATOM	210	C	ILE	A	42	19.161	31.290	11.832	1.00	23.98	C
ATOM	211	O	ILE	A	42	18.050	30.793	11.632	1.00	24.20	O
ATOM	212	CB	ILE	A	42	19.530	31.283	14.302	1.00	24.58	C
ATOM	213	CG1	ILE	A	42	19.779	32.181	15.518	1.00	22.71	C
ATOM	214	CG2	ILE	A	42	20.644	30.233	14.113	1.00	24.95	C
ATOM	215	CD1	ILE	A	42	19.466	31.516	16.837	1.00	22.70	C
ATOM	216	N	PRO	A	43	20.168	31.122	10.989	1.00	23.44	N
ATOM	217	CA	PRO	A	43	20.016	30.276	9.807	1.00	23.74	C
ATOM	218	C	PRO	A	43	19.709	28.802	10.154	1.00	24.82	C
ATOM	219	O	PRO	A	43	20.281	28.292	11.130	1.00	24.03	O
ATOM	220	CB	PRO	A	43	21.372	30.397	9.107	1.00	24.23	C
ATOM	221	CG	PRO	A	43	22.071	31.602	9.740	1.00	23.91	C
ATOM	222	CD	PRO	A	43	21.504	31.737	11.092	1.00	23.12	C
ATOM	223	N	ARG	A	44	18.784	28.178	9.406	1.00	25.20	N
ATOM	224	CA	ARG	A	44	18.439	26.767	9.520	1.00	26.19	C
ATOM	225	C	ARG	A	44	18.977	26.125	8.269	1.00	25.69	C
ATOM	226	O	ARG	A	44	18.563	26.468	7.159	1.00	25.67	O
ATOM	227	CB	ARG	A	44	16.934	26.504	9.513	1.00	27.16	C
ATOM	228	CG	ARG	A	44	16.140	27.062	10.678	1.00	31.24	C
ATOM	229	CD	ARG	A	44	14.653	26.529	10.769	1.00	32.98	C
ATOM	230	NE	ARG	A	44	14.400	25.143	10.311	1.00	34.09	C
ATOM	231	CZ	ARG	A	44	14.258	24.070	11.128	1.00	33.28	N
ATOM	232	NH1	ARG	A	44	14.384	24.177	12.453	1.00	29.63	C
ATOM	233	NH2	ARG	A	44	13.995	22.873	10.617	1.00	34.28	N
ATOM	234	N	LEU	A	45	19.870	25.174	8.433	1.00	24.68	N

ATOM	235	CA	LEU	A	45	20.551	24.608	7.302	1.00	24.30
ATOM	236	C	LEU	A	45	20.768	23.134	7.471	1.00	24.39
ATOM	237	O	LEU	A	45	20.711	22.623	8.588	1.00	23.97
ATOM	238	CB	LEU	A	45	21.934	25.233	7.205	1.00	23.42
ATOM	239	CG	LEU	A	45	21.929	26.724	6.941	1.00	24.95
ATOM	240	CD1	LEU	A	45	23.339	27.284	7.080	1.00	26.26
ATOM	241	CD2	LEU	A	45	21.375	26.953	5.533	1.00	24.54
ATOM	242	N	SER	A	46	21.092	22.494	6.352	1.00	24.71
ATOM	243	CA	SER	A	46	21.498	21.119	6.345	1.00	25.24
ATOM	244	C	SER	A	46	22.930	21.043	6.804	1.00	25.58
ATOM	245	O	SER	A	46	23.741	21.908	6.522	1.00	24.65
ATOM	246	CB	SER	A	46	21.401	20.505	4.950	1.00	25.11
ATOM	247	OG	SER	A	46	21.863	19.150	4.965	1.00	24.50
ATOM	248	N	GLN	A	47	23.208	19.970	7.517	1.00	26.68
ATOM	249	CA	GLN	A	47	24.524	19.626	8.011	1.00	27.63
ATOM	250	C	GLN	A	47	25.510	19.442	6.853	1.00	28.01
ATOM	251	O	GLN	A	47	26.704	19.613	7.026	1.00	28.15
ATOM	252	CB	GLN	A	47	24.368	18.317	8.803	1.00	28.89
ATOM	253	CG	GLN	A	47	25.580	17.447	8.920	1.00	31.28
ATOM	254	CD	GLN	A	47	25.826	16.556	7.765	1.00	32.33
ATOM	255	OE1	GLN	A	47	24.906	16.149	7.048	1.00	37.08
ATOM	256	NE2	GLN	A	47	27.089	16.213	7.576	1.00	35.81
ATOM	257	N	SER	A	48	25.026	19.086	5.667	1.00	28.11
ATOM	258	CA	SER	A	48	25.930	18.925	4.524	1.00	28.72
ATOM	259	C	SER	A	48	26.222	20.262	3.828	1.00	29.21
ATOM	260	O	SER	A	48	27.068	20.339	2.936	1.00	29.21
ATOM	261	CB	SER	A	48	25.343	17.952	3.508	1.00	28.66
ATOM	262	OG	SER	A	48	24.111	18.440	2.993	1.00	29.43
ATOM	263	N	ASP	A	49	25.525	21.314	4.236	1.00	29.60
ATOM	264	CA	ASP	A	49	25.683	22.610	3.596	1.00	30.12
ATOM	265	C	ASP	A	49	26.949	23.307	4.087	1.00	30.43
ATOM	266	O	ASP	A	49	27.100	23.565	5.272	1.00	29.60
ATOM	267	CB	ASP	A	49	24.450	23.458	3.858	1.00	30.26
ATOM	268	CG	ASP	A	49	24.491	24.792	3.151	1.00	31.74
ATOM	269	OD1	ASP	A	49	25.589	25.306	2.859	1.00	31.79
ATOM	270	OD2	ASP	A	49	23.454	25.418	2.878	1.00	34.13
ATOM	271	N	PRO	A	50	27.829	23.668	3.155	1.00	31.31
ATOM	272	CA	PRO	A	50	29.123	24.273	3.499	1.00	31.63
ATOM	273	C	PRO	A	50	28.965	25.514	4.355	1.00	31.72
ATOM	274	O	PRO	A	50	29.849	25.807	5.164	1.00	31.85
ATOM	275	CB	PRO	A	50	29.726	24.643	2.124	1.00	31.53
ATOM	276	CG	PRO	A	50	29.036	23.738	1.168	1.00	32.13
ATOM	277	CD	PRO	A	50	27.624	23.620	1.697	1.00	31.55
ATOM	278	N	ARG	A	51	27.875	26.246	4.171	1.00	32.04
ATOM	279	CA	ARG	A	51	27.648	27.443	4.962	1.00	32.56
ATOM	280	C	ARG	A	51	27.478	27.078	6.439	1.00	32.09
ATOM	281	O	ARG	A	51	27.853	27.855	7.322	1.00	31.87
ATOM	282	CB	ARG	A	51	26.420	28.221	4.464	1.00	32.95
ATOM	283	CG	ARG	A	51	26.568	28.810	3.064	1.00	34.79
ATOM	284	CD	ARG	A	51	25.273	29.372	2.480	1.00	35.89
ATOM	285	NE	ARG	A	51	24.276	28.325	2.228	1.00	37.56
ATOM	286	CZ	ARG	A	51	22.982	28.564	2.061	1.00	39.54
ATOM	287	NH1	ARG	A	51	22.531	29.814	2.124	1.00	40.85
ATOM	288	NH2	ARG	A	51	22.135	27.573	1.823	1.00	38.55
ATOM	289	N	ALA	A	52	26.932	25.903	6.721	1.00	31.50
ATOM	290	CA	ALA	A	52	26.769	25.513	8.117	1.00	31.34
ATOM	291	C	ALA	A	52	28.135	25.310	8.744	1.00	31.24
ATOM	292	O	ALA	A	52	28.409	25.763	9.865	1.00	30.91
ATOM	293	CB	ALA	A	52	25.953	24.284	8.236	1.00	31.35
ATOM	294	N	GLU	A	53	29.005	24.652	8.005	1.00	30.92
ATOM	295	CA	GLU	A	53	30.332	24.404	8.501	1.00	31.92

[illegible]

ATOM	296	C	GLU	A	53	31.059	25.730	8.728	1.00	31.52	C
ATOM	297	O	GLU	A	53	31.766	25.898	9.717	1.00	30.52	O
ATOM	298	CB	GLU	A	53	31.119	23.554	7.525	1.00	32.10	C
ATOM	299	CG	GLU	A	53	32.216	22.795	8.229	1.00	36.08	C
ATOM	300	CD	GLU	A	53	31.776	21.411	8.717	1.00	39.44	C
ATOM	301	OE1	GLU	A	53	30.629	21.234	9.167	1.00	40.34	O
ATOM	302	OE2	GLU	A	53	32.606	20.487	8.652	1.00	43.45	O
ATOM	303	N	GLU	A	54	30.870	26.665	7.808	1.00	31.32	N
ATOM	304	CA	GLU	A	54	31.507	27.968	7.919	1.00	32.09	C
ATOM	305	C	GLU	A	54	31.039	28.685	9.193	1.00	31.07	C
ATOM	306	O	GLU	A	54	31.833	29.295	9.881	1.00	31.27	O
ATOM	307	CB	GLU	A	54	31.218	28.812	6.681	1.00	32.42	C
ATOM	308	CG	GLU	A	54	31.939	30.146	6.669	1.00	37.01	C
ATOM	309	CD	GLU	A	54	31.662	30.966	5.410	1.00	40.45	C
ATOM	310	OE1	GLU	A	54	30.843	30.528	4.558	1.00	41.94	O
ATOM	311	OE2	GLU	A	54	32.268	32.051	5.285	1.00	43.43	O
ATOM	312	N	LEU	A	55	29.761	28.567	9.534	1.00	29.85	N
ATOM	313	CA	LEU	A	55	29.251	29.240	10.708	1.00	28.58	C
ATOM	314	C	LEU	A	55	29.848	28.667	11.983	1.00	27.91	C
ATOM	315	O	LEU	A	55	30.304	29.419	12.841	1.00	26.43	O
ATOM	316	CB	LEU	A	55	27.734	29.179	10.733	1.00	28.61	C
ATOM	317	CG	LEU	A	55	27.097	30.041	9.652	1.00	28.69	C
ATOM	318	CD1	LEU	A	55	25.647	29.633	9.423	1.00	29.69	C
ATOM	319	CD2	LEU	A	55	27.193	31.507	10.005	1.00	27.60	C
ATOM	320	N	ILE	A	56	29.889	27.334	12.086	1.00	27.51	N
ATOM	321	CA	ILE	A	56	30.432	26.679	13.278	1.00	27.49	C
ATOM	322	C	ILE	A	56	31.910	27.014	13.457	1.00	28.32	C
ATOM	323	O	ILE	A	56	32.359	27.353	14.549	1.00	28.51	O
ATOM	324	CB	ILE	A	56	30.260	25.174	13.192	1.00	27.21	C
ATOM	325	CG1	ILE	A	56	28.771	24.787	13.218	1.00	25.69	C
ATOM	326	CG2	ILE	A	56	30.982	24.505	14.354	1.00	27.65	C
ATOM	327	CD1	ILE	A	56	28.484	23.341	12.760	1.00	23.98	C
ATOM	328	N	GLU	A	57	32.645	26.935	12.357	1.00	28.85	N
ATOM	329	CA	GLU	A	57	34.068	27.230	12.308	1.00	29.98	C
ATOM	330	C	GLU	A	57	34.319	28.623	12.837	1.00	30.12	C
ATOM	331	O	GLU	A	57	35.306	28.869	13.524	1.00	31.00	O
ATOM	332	CB	GLU	A	57	34.557	27.137	10.850	1.00	30.52	C
ATOM	333	CG	GLU	A	57	36.003	27.518	10.610	1.00	33.37	C
ATOM	334	CD	GLU	A	57	36.968	26.742	11.482	1.00	38.19	C
ATOM	335	OE1	GLU	A	57	36.689	25.558	11.810	1.00	41.02	O
ATOM	336	OE2	GLU	A	57	38.015	27.329	11.847	1.00	42.74	O
ATOM	337	N	ASN	A	58	33.409	29.527	12.516	1.00	29.57	N
ATOM	338	CA	ASN	A	58	33.516	30.911	12.929	1.00	29.62	C
ATOM	339	C	ASN	A	58	32.852	31.222	14.230	1.00	27.88	C
ATOM	340	O	ASN	A	58	32.690	32.364	14.566	1.00	26.74	O
ATOM	341	CB	ASN	A	58	32.805	31.769	11.918	1.00	30.43	C
ATOM	342	CG	ASN	A	58	33.719	32.439	11.040	1.00	33.87	C
ATOM	343	OD1	ASN	A	58	34.040	31.918	9.965	1.00	36.67	O
ATOM	344	ND2	ASN	A	58	34.189	33.629	11.467	1.00	38.66	N
ATOM	345	N	GLU	A	59	32.396	30.205	14.922	1.00	27.73	N
ATOM	346	CA	GLU	A	59	31.753	30.411	16.205	1.00	27.50	C
ATOM	347	C	GLU	A	59	30.545	31.320	16.104	1.00	26.07	C
ATOM	348	O	GLU	A	59	30.366	32.255	16.861	1.00	25.28	O
ATOM	349	CB	GLU	A	59	32.795	30.867	17.224	1.00	28.12	C
ATOM	350	CG	GLU	A	59	33.720	29.688	17.531	1.00	31.06	C
ATOM	351	CD	GLU	A	59	34.739	29.965	18.604	1.00	35.24	C
ATOM	352	OE1	GLU	A	59	35.826	30.432	18.239	1.00	39.24	O
ATOM	353	OE2	GLU	A	59	34.469	29.698	19.795	1.00	37.54	O
ATOM	354	N	GLU	A	60	29.691	30.982	15.155	1.00	25.70	N
ATOM	355	CA	GLU	A	60	28.417	31.643	14.971	1.00	25.50	C
ATOM	356	C	GLU	A	60	27.345	30.576	14.998	1.00	24.10	C

ATOM	357	O	GLU	A	60	27.527	29.479	14.487	1.00	22.92
ATOM	358	CB	GLU	A	60	28.402	32.404	13.664	1.00	25.96
ATOM	359	CG	GLU	A	60	29.454	33.497	13.697	1.00	30.39
ATOM	360	CD	GLU	A	60	29.218	34.599	12.703	1.00	35.90
ATOM	361	OE1	GLU	A	60	29.281	34.349	11.478	1.00	38.45
ATOM	362	OE2	GLU	A	60	28.998	35.732	13.165	1.00	43.84
ATOM	363	N	PRO	A	61	26.203	30.933	15.546	1.00	22.92
ATOM	364	CA	PRO	A	61	25.112	29.983	15.735	1.00	22.31
ATOM	365	C	PRO	A	61	24.482	29.519	14.445	1.00	21.76
ATOM	366	O	PRO	A	61	24.394	30.252	13.474	1.00	21.59
ATOM	367	CB	PRO	A	61	24.079	30.775	16.525	1.00	22.55
ATOM	368	CG	PRO	A	61	24.450	32.218	16.380	1.00	21.42
ATOM	369	CD	PRO	A	61	25.863	32.295	15.967	1.00	22.38
ATOM	370	N	VAL	A	62	24.032	28.279	14.444	1.00	21.55
ATOM	371	CA	VAL	A	62	23.306	27.739	13.317	1.00	21.90
ATOM	372	C	VAL	A	62	22.415	26.622	13.817	1.00	22.38
ATOM	373	O	VAL	A	62	22.771	25.886	14.757	1.00	22.77
ATOM	374	CB	VAL	A	62	24.262	27.217	12.218	1.00	22.10
ATOM	375	CG1	VAL	A	62	25.134	26.112	12.738	1.00	22.89
ATOM	376	CG2	VAL	A	62	23.486	26.748	11.000	1.00	21.98
ATOM	377	N	VAL	A	63	21.234	26.517	13.226	1.00	22.90
ATOM	378	CA	VAL	A	63	20.373	25.398	13.499	1.00	23.20
ATOM	379	C	VAL	A	63	20.586	24.373	12.378	1.00	23.18
ATOM	380	O	VAL	A	63	20.395	24.682	11.208	1.00	23.38
ATOM	381	CB	VAL	A	63	18.880	25.768	13.561	1.00	23.27
ATOM	382	CG1	VAL	A	63	18.046	24.483	13.754	1.00	24.19
ATOM	383	CG2	VAL	A	63	18.620	26.695	14.692	1.00	22.51
ATOM	384	N	LEU	A	64	21.007	23.181	12.756	1.00	23.15
ATOM	385	CA	LEU	A	64	21.158	22.060	11.835	1.00	24.29
ATOM	386	C	LEU	A	64	19.904	21.203	11.906	1.00	23.48
ATOM	387	O	LEU	A	64	19.488	20.816	12.988	1.00	23.20
ATOM	388	CB	LEU	A	64	22.383	21.235	12.206	1.00	24.79
ATOM	389	CG	LEU	A	64	23.649	22.089	12.141	1.00	28.22
ATOM	390	CD1	LEU	A	64	24.810	21.338	12.699	1.00	32.80
ATOM	391	CD2	LEU	A	64	23.946	22.478	10.719	1.00	30.04
ATOM	392	N	THR	A	65	19.308	20.910	10.759	1.00	22.79
ATOM	393	CA	THR	A	65	18.019	20.218	10.740	1.00	23.27
ATOM	394	C	THR	A	65	18.088	18.730	10.534	1.00	23.05
ATOM	395	O	THR	A	65	17.102	18.050	10.765	1.00	22.85
ATOM	396	CB	THR	A	65	17.150	20.724	9.571	1.00	23.53
ATOM	397	OG1	THR	A	65	17.855	20.496	8.352	1.00	22.75
ATOM	398	CG2	THR	A	65	16.966	22.226	9.609	1.00	24.78
ATOM	399	N	ASP	A	66	19.231	18.227	10.093	1.00	23.33
ATOM	400	CA	ASP	A	66	19.330	16.828	9.731	1.00	23.76
ATOM	401	C	ASP	A	66	20.581	16.076	10.209	1.00	23.41
ATOM	402	O	ASP	A	66	21.117	15.265	9.457	1.00	23.64
ATOM	403	CB	ASP	A	66	19.245	16.732	8.199	1.00	23.97
ATOM	404	CG	ASP	A	66	20.326	17.517	7.514	1.00	24.64
ATOM	405	OD1	ASP	A	66	21.175	18.099	8.223	1.00	23.06
ATOM	406	OD2	ASP	A	66	20.417	17.612	6.268	1.00	26.54
ATOM	407	N	THR	A	67	21.050	16.336	11.428	1.00	22.72
ATOM	408	CA	THR	A	67	22.235	15.655	11.912	1.00	21.85
ATOM	409	C	THR	A	67	21.928	14.246	12.349	1.00	21.26
ATOM	410	O	THR	A	67	22.813	13.429	12.353	1.00	20.41
ATOM	411	CB	THR	A	67	22.836	16.345	13.138	1.00	21.92
ATOM	412	OG1	THR	A	67	21.884	16.347	14.200	1.00	20.59
ATOM	413	CG2	THR	A	67	23.161	17.822	12.873	1.00	23.29
ATOM	414	N	ASN	A	68	20.684	13.993	12.752	1.00	20.82
ATOM	415	CA	ASN	A	68	20.321	12.741	13.386	1.00	21.01
ATOM	416	C	ASN	A	68	21.146	12.525	14.652	1.00	20.47
ATOM	417	O	ASN	A	68	21.370	11.377	15.069	1.00	20.06

O
C
C
C
O
O
N
C
C
O
C
C
N
C
C
O
C
C
N
C
C
O
C
C
C
N
C
C
O
C
C
C
N
C
C
O
C
C
O
N
C
C
O
C
N
C
C
O

ATOM	418	CB	ASN	A	68	20.516	11.546	12.444	1.00	21.63	C
ATOM	419	CG	ASN	A	68	19.476	11.493	11.340	1.00	22.28	C
ATOM	420	OD1	ASN	A	68	18.276	11.429	11.601	1.00	23.83	O
ATOM	421	ND2	ASN	A	68	19.936	11.492	10.108	1.00	22.82	N
ATOM	422	N	LEU	A	69	21.613	13.617	15.248	1.00	19.86	N
ATOM	423	CA	LEU	A	69	22.442	13.533	16.464	1.00	19.68	C
ATOM	424	C	LEU	A	69	21.814	12.707	17.571	1.00	19.03	C
ATOM	425	O	LEU	A	69	22.492	11.878	18.169	1.00	18.33	O
ATOM	426	CB	LEU	A	69	22.770	14.907	17.009	1.00	19.89	C
ATOM	427	CG	LEU	A	69	23.654	14.999	18.239	1.00	19.87	C
ATOM	428	CD1	LEU	A	69	24.970	14.289	18.035	1.00	20.74	C
ATOM	429	CD2	LEU	A	69	23.911	16.459	18.577	1.00	20.41	C
ATOM	430	N	VAL	A	70	20.538	12.940	17.860	1.00	18.84	N
ATOM	431	CA	VAL	A	70	19.843	12.177	18.894	1.00	18.85	C
ATOM	432	C	VAL	A	70	18.634	11.447	18.339	1.00	19.17	C
ATOM	433	O	VAL	A	70	17.604	11.325	19.000	1.00	18.71	O
ATOM	434	CB	VAL	A	70	19.418	13.038	20.097	1.00	18.97	C
ATOM	435	CG1	VAL	A	70	20.645	13.557	20.812	1.00	20.41	C
ATOM	436	CG2	VAL	A	70	18.513	14.185	19.686	1.00	18.87	C
ATOM	437	N	TYR	A	71	18.796	10.916	17.133	1.00	20.12	N
ATOM	438	CA	TYR	A	71	17.711	10.224	16.454	1.00	20.64	C
ATOM	439	C	TYR	A	71	17.003	9.218	17.377	1.00	20.76	C
ATOM	440	O	TYR	A	71	15.804	9.275	17.507	1.00	20.54	O
ATOM	441	CB	TYR	A	71	18.186	9.591	15.136	1.00	20.75	C
ATOM	442	CG	TYR	A	71	17.243	8.517	14.628	1.00	23.02	C
ATOM	443	CD1	TYR	A	71	16.012	8.837	14.046	1.00	24.62	C
ATOM	444	CD2	TYR	A	71	17.572	7.182	14.754	1.00	22.94	C
ATOM	445	CE1	TYR	A	71	15.132	7.800	13.595	1.00	23.64	C
ATOM	446	CE2	TYR	A	71	16.730	6.174	14.312	1.00	23.48	C
ATOM	447	CZ	TYR	A	71	15.524	6.478	13.730	1.00	24.03	C
ATOM	448	OH	TYR	A	71	14.697	5.422	13.349	1.00	29.55	O
ATOM	449	N	PRO	A	72	17.728	8.356	18.068	1.00	21.35	N
ATOM	450	CA	PRO	A	72	17.080	7.363	18.945	1.00	21.89	C
ATOM	451	C	PRO	A	72	16.296	7.955	20.106	1.00	22.22	C
ATOM	452	O	PRO	A	72	15.432	7.269	20.628	1.00	21.25	O
ATOM	453	CB	PRO	A	72	18.248	6.538	19.493	1.00	21.83	C
ATOM	454	CG	PRO	A	72	19.420	6.849	18.606	1.00	22.53	C
ATOM	455	CD	PRO	A	72	19.192	8.240	18.080	1.00	21.81	C
ATOM	456	N	ALA	A	73	16.568	9.206	20.484	1.00	22.77	N
ATOM	457	CA	ALA	A	73	15.859	9.836	21.605	1.00	23.23	C
ATOM	458	C	ALA	A	73	14.542	10.487	21.178	1.00	23.64	C
ATOM	459	O	ALA	A	73	13.764	10.966	22.014	1.00	23.47	O
ATOM	460	CB	ALA	A	73	16.750	10.860	22.284	1.00	22.77	C
ATOM	461	N	LEU	A	74	14.275	10.507	19.881	1.00	24.25	N
ATOM	462	CA	LEU	A	74	13.057	11.148	19.405	1.00	24.86	C
ATOM	463	C	LEU	A	74	11.792	10.466	19.920	1.00	25.46	C
ATOM	464	O	LEU	A	74	10.729	11.070	19.967	1.00	25.11	O
ATOM	465	CB	LEU	A	74	13.055	11.240	17.881	1.00	24.77	C
ATOM	466	CG	LEU	A	74	14.160	12.161	17.338	1.00	26.74	C
ATOM	467	CD1	LEU	A	74	13.915	12.478	15.867	1.00	28.86	C
ATOM	468	CD2	LEU	A	74	14.288	13.447	18.135	1.00	26.86	C
ATOM	469	N	LYS	A	75	11.908	9.204	20.306	1.00	26.12	N
ATOM	470	CA	LYS	A	75	10.756	8.460	20.814	1.00	26.39	C
ATOM	471	C	LYS	A	75	10.585	8.687	22.309	1.00	26.71	C
ATOM	472	O	LYS	A	75	9.584	8.284	22.875	1.00	26.80	O
ATOM	473	CB	LYS	A	75	10.918	6.963	20.534	1.00	26.36	C
ATOM	474	CG	LYS	A	75	12.141	6.322	21.185	1.00	26.49	C
ATOM	475	CD	LYS	A	75	12.254	4.813	20.838	1.00	27.59	C
ATOM	476	CE	LYS	A	75	13.732	4.367	20.671	1.00	27.21	C
ATOM	477	NZ	LYS	A	75	14.475	4.226	21.912	1.00	24.71	N
ATOM	478	N	TRP	A	76	11.561	9.327	22.950	1.00	26.64	N

ATOM	479	CA	TRP	A	76	11.484	9.557	24.383	1.00	26.78
ATOM	480	C	TRP	A	76	10.253	10.368	24.770	1.00	27.48
ATOM	481	O	TRP	A	76	9.889	11.317	24.095	1.00	27.93
ATOM	482	CB	TRP	A	76	12.717	10.311	24.888	1.00	26.57
ATOM	483	CG	TRP	A	76	13.963	9.500	24.895	1.00	25.80
ATOM	484	CD1	TRP	A	76	14.101	8.219	24.486	1.00	24.26
ATOM	485	CD2	TRP	A	76	15.255	9.917	25.347	1.00	22.71
ATOM	486	NE1	TRP	A	76	15.399	7.804	24.647	1.00	23.20
ATOM	487	CE2	TRP	A	76	16.128	8.829	25.178	1.00	22.53
ATOM	488	CE3	TRP	A	76	15.767	11.107	25.867	1.00	21.96
ATOM	489	CZ2	TRP	A	76	17.468	8.890	25.520	1.00	22.50
ATOM	490	CZ3	TRP	A	76	17.090	11.172	26.202	1.00	20.53
ATOM	491	CH2	TRP	A	76	17.931	10.076	26.029	1.00	22.36
ATOM	492	N	ASP	A	77	9.639	9.976	25.880	1.00	27.85
ATOM	493	CA	ASP	A	77	8.532	10.684	26.484	1.00	27.74
ATOM	494	C	ASP	A	77	8.560	10.277	27.957	1.00	27.45
ATOM	495	O	ASP	A	77	9.373	9.460	28.336	1.00	27.22
ATOM	496	CB	ASP	A	77	7.208	10.368	25.800	1.00	27.89
ATOM	497	CG	ASP	A	77	6.802	8.913	25.903	1.00	29.06
ATOM	498	OD1	ASP	A	77	7.354	8.108	26.708	1.00	30.51
ATOM	499	OD2	ASP	A	77	5.894	8.489	25.179	1.00	31.13
ATOM	500	N	LEU	A	78	7.710	10.847	28.796	1.00	27.47
ATOM	501	CA	LEU	A	78	7.819	10.578	30.229	1.00	27.63
ATOM	502	C	LEU	A	78	7.655	9.101	30.542	1.00	27.81
ATOM	503	O	LEU	A	78	8.386	8.548	31.367	1.00	27.16
ATOM	504	CB	LEU	A	78	6.818	11.409	31.011	1.00	27.59
ATOM	505	CG	LEU	A	78	7.007	12.916	30.880	1.00	28.68
ATOM	506	CD1	LEU	A	78	5.906	13.677	31.612	1.00	28.95
ATOM	507	CD2	LEU	A	78	8.369	13.328	31.412	1.00	29.32
ATOM	508	N	GLU	A	79	6.721	8.450	29.860	1.00	28.17
ATOM	509	CA	GLU	A	79	6.480	7.035	30.102	1.00	28.56
ATOM	510	C	GLU	A	79	7.715	6.185	29.781	1.00	28.22
ATOM	511	O	GLU	A	79	8.167	5.388	30.600	1.00	27.73
ATOM	512	CB	GLU	A	79	5.267	6.540	29.298	1.00	28.87
ATOM	513	CG	GLU	A	79	5.051	5.049	29.467	1.00	31.45
ATOM	514	CD	GLU	A	79	3.849	4.516	28.716	1.00	34.79
ATOM	515	OE1	GLU	A	79	3.422	5.129	27.709	1.00	35.78
ATOM	516	OE2	GLU	A	79	3.343	3.457	29.144	1.00	37.92
ATOM	517	N	TYR	A	80	8.260	6.346	28.582	1.00	28.27
ATOM	518	CA	TYR	A	80	9.452	5.597	28.200	1.00	28.00
ATOM	519	C	TYR	A	80	10.628	5.873	29.142	1.00	27.64
ATOM	520	O	TYR	A	80	11.330	4.958	29.563	1.00	27.16
ATOM	521	CB	TYR	A	80	9.838	5.964	26.787	1.00	28.13
ATOM	522	CG	TYR	A	80	11.054	5.250	26.246	1.00	27.95
ATOM	523	CD1	TYR	A	80	10.952	3.991	25.644	1.00	27.39
ATOM	524	CD2	TYR	A	80	12.300	5.846	26.306	1.00	25.55
ATOM	525	CE1	TYR	A	80	12.089	3.352	25.128	1.00	26.82
ATOM	526	CE2	TYR	A	80	13.417	5.224	25.808	1.00	25.50
ATOM	527	CZ	TYR	A	80	13.320	3.992	25.213	1.00	25.64
ATOM	528	OH	TYR	A	80	14.467	3.426	24.703	1.00	23.50
ATOM	529	N	LEU	A	81	10.839	7.135	29.482	1.00	27.51
ATOM	530	CA	LEU	A	81	11.952	7.477	30.371	1.00	27.75
ATOM	531	C	LEU	A	81	11.741	6.943	31.801	1.00	27.76
ATOM	532	O	LEU	A	81	12.682	6.459	32.437	1.00	27.35
ATOM	533	CB	LEU	A	81	12.194	8.990	30.399	1.00	27.35
ATOM	534	CG	LEU	A	81	12.659	9.656	29.096	1.00	28.06
ATOM	535	CD1	LEU	A	81	12.664	11.170	29.269	1.00	28.55
ATOM	536	CD2	LEU	A	81	14.036	9.192	28.655	1.00	27.65
ATOM	537	N	GLN	A	82	10.521	7.049	32.313	1.00	28.10
ATOM	538	CA	GLN	A	82	10.234	6.557	33.656	1.00	28.61
ATOM	539	C	GLN	A	82	10.525	5.066	33.689	1.00	28.32

[illegible]

ATOM	540	O	GLN	A	82	11.070	4.550	34.643	1.00	28.25	O
ATOM	541	CB	GLN	A	82	8.774	6.805	34.032	1.00	29.04	C
ATOM	542	CG	GLN	A	82	8.325	6.063	35.293	1.00	30.18	C
ATOM	543	CD	GLN	A	82	7.184	6.754	36.042	1.00	32.25	C
ATOM	544	OE1	GLN	A	82	6.642	7.758	35.594	1.00	34.51	O
ATOM	545	NE2	GLN	A	82	6.822	6.204	37.189	1.00	36.79	N
ATOM	546	N	GLU	A	83	10.193	4.397	32.601	1.00	28.58	N
ATOM	547	CA	GLU	A	83	10.349	2.950	32.492	1.00	28.70	C
ATOM	548	C	GLU	A	83	11.801	2.515	32.314	1.00	28.14	C
ATOM	549	O	GLU	A	83	12.166	1.424	32.713	1.00	27.54	O
ATOM	550	CB	GLU	A	83	9.506	2.438	31.319	1.00	28.53	C
ATOM	551	CG	GLU	A	83	9.562	0.937	31.101	1.00	30.76	C
ATOM	552	CD	GLU	A	83	8.985	0.150	32.265	1.00	33.31	C
ATOM	553	OE1	GLU	A	83	8.172	0.713	33.030	1.00	35.22	O
ATOM	554	OE2	GLU	A	83	9.352	-1.033	32.428	1.00	35.98	O
ATOM	555	N	ASN	A	84	12.644	3.374	31.753	1.00	27.44	N
ATOM	556	CA	ASN	A	84	13.985	2.938	31.400	1.00	27.09	C
ATOM	557	C	ASN	A	84	15.168	3.729	31.913	1.00	27.31	C
ATOM	558	O	ASN	A	84	16.291	3.276	31.759	1.00	27.17	O
ATOM	559	CB	ASN	A	84	14.099	2.917	29.879	1.00	27.17	C
ATOM	560	CG	ASN	A	84	13.226	1.890	29.254	1.00	26.24	C
ATOM	561	OD1	ASN	A	84	13.361	0.696	29.532	1.00	27.49	O
ATOM	562	ND2	ASN	A	84	12.312	2.333	28.413	1.00	23.37	N
ATOM	563	N	ILE	A	85	14.952	4.893	32.511	1.00	27.52	N
ATOM	564	CA	ILE	A	85	16.088	5.734	32.861	1.00	28.25	C
ATOM	565	C	ILE	A	85	16.788	5.390	34.185	1.00	28.03	C
ATOM	566	O	ILE	A	85	17.700	6.094	34.610	1.00	28.40	O
ATOM	567	CB	ILE	A	85	15.684	7.221	32.801	1.00	28.23	C
ATOM	568	CG1	ILE	A	85	16.872	8.069	32.342	1.00	29.61	C
ATOM	569	CG2	ILE	A	85	15.143	7.694	34.139	1.00	29.11	C
ATOM	570	CD1	ILE	A	85	16.520	9.535	31.996	1.00	28.97	C
ATOM	571	N	GLY	A	86	16.368	4.317	34.833	1.00	28.14	N
ATOM	572	CA	GLY	A	86	17.014	3.874	36.061	1.00	28.15	C
ATOM	573	C	GLY	A	86	16.478	4.504	37.335	1.00	28.11	C
ATOM	574	O	GLY	A	86	15.494	5.251	37.308	1.00	27.93	O
ATOM	575	N	ASN	A	87	17.162	4.220	38.444	1.00	27.96	N
ATOM	576	CA	ASN	A	87	16.754	4.672	39.767	1.00	27.88	C
ATOM	577	C	ASN	A	87	17.736	5.677	40.373	1.00	27.82	C
ATOM	578	O	ASN	A	87	17.751	5.886	41.585	1.00	27.63	O
ATOM	579	CB	ASN	A	87	16.571	3.447	40.713	1.00	27.89	C
ATOM	580	N	GLY	A	88	18.559	6.301	39.538	1.00	28.21	N
ATOM	581	CA	GLY	A	88	19.478	7.336	40.006	1.00	28.18	C
ATOM	582	C	GLY	A	88	18.706	8.584	40.396	1.00	28.19	C
ATOM	583	O	GLY	A	88	17.520	8.676	40.097	1.00	28.62	O
ATOM	584	N	ASP	A	89	19.357	9.530	41.068	1.00	27.99	N
ATOM	585	CA	ASP	A	89	18.707	10.781	41.468	1.00	28.09	C
ATOM	586	C	ASP	A	89	18.655	11.806	40.335	1.00	28.00	C
ATOM	587	O	ASP	A	89	19.557	11.866	39.507	1.00	28.07	O
ATOM	588	CB	ASP	A	89	19.467	11.433	42.613	1.00	28.24	C
ATOM	589	CG	ASP	A	89	19.249	10.747	43.935	1.00	28.53	C
ATOM	590	OD1	ASP	A	89	18.398	9.843	44.024	1.00	30.26	O
ATOM	591	OD2	ASP	A	89	19.884	11.070	44.955	1.00	29.31	O
ATOM	592	N	PHE	A	90	17.620	12.637	40.330	1.00	27.56	N
ATOM	593	CA	PHE	A	90	17.510	13.687	39.332	1.00	27.37	C
ATOM	594	C	PHE	A	90	17.291	15.037	39.993	1.00	27.78	C
ATOM	595	O	PHE	A	90	16.453	15.171	40.884	1.00	27.69	O
ATOM	596	CB	PHE	A	90	16.378	13.388	38.351	1.00	27.05	C
ATOM	597	CG	PHE	A	90	16.678	12.262	37.408	1.00	25.85	C
ATOM	598	CD1	PHE	A	90	16.525	10.943	37.806	1.00	25.70	C
ATOM	599	CD2	PHE	A	90	17.119	12.521	36.121	1.00	26.20	C
ATOM	600	CE1	PHE	A	90	16.797	9.908	36.943	1.00	24.06	C

ATOM	601	CE2	PHE	A	90	17.395	11.489	35.248	1.00	25.49
ATOM	602	CZ	PHE	A	90	17.235	10.176	35.666	1.00	25.33
ATOM	603	N	SER	A	91	18.062	16.033	39.560	1.00	27.85
ATOM	604	CA	SER	A	91	17.902	17.384	40.075	1.00	28.17
ATOM	605	C	SER	A	91	16.695	18.023	39.419	1.00	28.48
ATOM	606	O	SER	A	91	16.589	18.085	38.192	1.00	28.35
ATOM	607	CB	SER	A	91	19.149	18.236	39.838	1.00	27.86
ATOM	608	OG	SER	A	91	20.260	17.652	40.485	1.00	27.15
ATOM	609	N	VAL	A	92	15.784	18.482	40.261	1.00	29.05
ATOM	610	CA	VAL	A	92	14.585	19.141	39.812	1.00	29.67
ATOM	611	C	VAL	A	92	14.437	20.465	40.529	1.00	30.31
ATOM	612	O	VAL	A	92	14.442	20.537	41.763	1.00	30.37
ATOM	613	CB	VAL	A	92	13.352	18.313	40.106	1.00	29.81
ATOM	614	CG1	VAL	A	92	12.106	19.026	39.582	1.00	29.93
ATOM	615	CG2	VAL	A	92	13.490	16.930	39.507	1.00	29.60
ATOM	616	N	TYR	A	93	14.329	21.516	39.732	1.00	31.00
ATOM	617	CA	TYR	A	93	14.135	22.845	40.241	1.00	31.29
ATOM	618	C	TYR	A	93	12.662	23.113	40.298	1.00	31.54
ATOM	619	O	TYR	A	93	11.892	22.701	39.423	1.00	31.35
ATOM	620	CB	TYR	A	93	14.829	23.853	39.345	1.00	31.87
ATOM	621	CG	TYR	A	93	16.310	23.711	39.436	1.00	33.05
ATOM	622	CD1	TYR	A	93	17.012	24.283	40.482	1.00	34.67
ATOM	623	CD2	TYR	A	93	16.998	22.954	38.525	1.00	34.24
ATOM	624	CE1	TYR	A	93	18.357	24.125	40.593	1.00	35.09
ATOM	625	CE2	TYR	A	93	18.338	22.789	38.634	1.00	36.36
ATOM	626	CZ	TYR	A	93	19.011	23.379	39.670	1.00	36.33
ATOM	627	OH	TYR	A	93	20.357	23.204	39.772	1.00	40.81
ATOM	628	N	SER	A	94	12.286	23.813	41.351	1.00	32.03
ATOM	629	CA	SER	A	94	10.919	24.182	41.599	1.00	32.31
ATOM	630	C	SER	A	94	10.884	25.692	41.704	1.00	32.42
ATOM	631	O	SER	A	94	11.743	26.286	42.350	1.00	32.48
ATOM	632	CB	SER	A	94	10.464	23.566	42.914	1.00	32.38
ATOM	633	OG	SER	A	94	9.062	23.691	43.052	1.00	33.53
ATOM	634	N	ALA	A	95	9.902	26.314	41.070	1.00	32.58
ATOM	635	CA	ALA	A	95	9.766	27.757	41.135	1.00	32.90
ATOM	636	C	ALA	A	95	8.312	28.172	41.076	1.00	33.48
ATOM	637	O	ALA	A	95	7.446	27.429	40.620	1.00	33.62
ATOM	638	CB	ALA	A	95	10.528	28.413	39.996	1.00	32.80
ATOM	639	N	SER	A	96	8.053	29.382	41.539	1.00	34.08
ATOM	640	CA	SER	A	96	6.721	29.932	41.482	1.00	34.75
ATOM	641	C	SER	A	96	6.616	30.974	40.382	1.00	34.55
ATOM	642	O	SER	A	96	5.603	31.659	40.275	1.00	35.29
ATOM	643	CB	SER	A	96	6.363	30.592	42.801	1.00	35.05
ATOM	644	OG	SER	A	96	5.165	31.311	42.627	1.00	36.18
ATOM	645	N	THR	A	97	7.673	31.107	39.593	1.00	33.88
ATOM	646	CA	THR	A	97	7.716	32.046	38.477	1.00	33.41
ATOM	647	C	THR	A	97	8.084	31.238	37.265	1.00	32.32
ATOM	648	O	THR	A	97	8.590	30.143	37.411	1.00	32.10
ATOM	649	CB	THR	A	97	8.797	33.144	38.695	1.00	33.64
ATOM	650	OG1	THR	A	97	9.067	33.810	37.460	1.00	33.76
ATOM	651	CG2	THR	A	97	10.190	32.559	39.058	1.00	34.25
ATOM	652	N	HIS	A	98	7.840	31.762	36.073	1.00	31.47
ATOM	653	CA	HIS	A	98	8.278	31.071	34.863	1.00	30.97
ATOM	654	C	HIS	A	98	9.804	31.134	34.707	1.00	30.53
ATOM	655	O	HIS	A	98	10.378	30.379	33.940	1.00	29.14
ATOM	656	CB	HIS	A	98	7.613	31.666	33.621	1.00	31.14
ATOM	657	CG	HIS	A	98	7.878	33.125	33.415	1.00	30.86
ATOM	658	ND1	HIS	A	98	7.121	34.112	34.011	1.00	31.51
ATOM	659	CD2	HIS	A	98	8.795	33.767	32.651	1.00	31.48
ATOM	660	CE1	HIS	A	98	7.566	35.298	33.634	1.00	31.14
ATOM	661	NE2	HIS	A	98	8.581	35.118	32.805	1.00	30.50

C
C
N
C
C
C
O
C
C
O
N
C
C
C
O
C
C
C
N
C
C
C
O
C
C
O
N
C
C
C
O
N
C
C
C
O
C
C
O
N
C
C
C
O
C
C
N
C
C
C
O
C
C
N
C
C
C
N

ATOM	662	N	LYS	A	99	10.459	32.025	35.449	1.00	30.36	
ATOM	663	CA	LYS	A	99	11.895	32.198	35.298	1.00	31.07	N
ATOM	664	C	LYS	A	99	12.761	31.288	36.166	1.00	31.00	C
ATOM	665	O	LYS	A	99	12.693	31.337	37.396	1.00	31.91	C
ATOM	666	CB	LYS	A	99	12.265	33.643	35.544	1.00	31.15	O
ATOM	667	CG	LYS	A	99	11.887	34.551	34.391	1.00	33.39	C
ATOM	668	CD	LYS	A	99	12.486	35.945	34.556	1.00	36.16	C
ATOM	669	CE	LYS	A	99	11.763	36.771	35.607	1.00	37.52	C
ATOM	670	NZ	LYS	A	99	10.620	37.507	35.006	1.00	38.84	C
ATOM	671	N	PHE	A	100	13.572	30.453	35.518	1.00	30.20	N
ATOM	672	CA	PHE	A	100	14.517	29.606	36.232	1.00	30.03	N
ATOM	673	C	PHE	A	100	15.952	30.123	36.138	1.00	29.72	C
ATOM	674	O	PHE	A	100	16.826	29.544	35.481	1.00	29.39	C
ATOM	675	CB	PHE	A	100	14.441	28.168	35.744	1.00	29.93	O
ATOM	676	CG	PHE	A	100	13.223	27.452	36.206	1.00	29.72	C
ATOM	677	CD1	PHE	A	100	12.014	27.637	35.566	1.00	28.98	C
ATOM	678	CD2	PHE	A	100	13.283	26.596	37.289	1.00	30.77	C
ATOM	679	CE1	PHE	A	100	10.896	26.969	35.983	1.00	30.22	C
ATOM	680	CE2	PHE	A	100	12.158	25.919	37.719	1.00	30.51	C
ATOM	681	CZ	PHE	A	100	10.967	26.105	37.069	1.00	30.63	C
ATOM	682	N	LEU	A	101	16.178	31.235	36.809	1.00	29.87	C
ATOM	683	CA	LEU	A	101	17.495	31.812	36.919	1.00	29.57	N
ATOM	684	C	LEU	A	101	18.451	30.814	37.555	1.00	29.61	C
ATOM	685	O	LEU	A	101	18.249	30.380	38.679	1.00	28.35	C
ATOM	686	CB	LEU	A	101	17.412	33.057	37.787	1.00	29.68	O
ATOM	687	CG	LEU	A	101	18.707	33.845	37.954	1.00	29.68	C
ATOM	688	CD1	LEU	A	101	19.184	34.337	36.620	1.00	28.70	C
ATOM	689	CD2	LEU	A	101	18.474	35.014	38.909	1.00	31.41	C
ATOM	690	N	TYR	A	102	19.490	30.440	36.816	1.00	30.36	C
ATOM	691	CA	TYR	A	102	20.516	29.535	37.338	1.00	30.64	N
ATOM	692	C	TYR	A	102	21.332	30.228	38.440	1.00	30.93	C
ATOM	693	O	TYR	A	102	21.623	31.425	38.339	1.00	30.67	C
ATOM	694	CB	TYR	A	102	21.480	29.101	36.224	1.00	30.45	O
ATOM	695	CG	TYR	A	102	22.609	28.271	36.774	1.00	31.25	C
ATOM	696	CD1	TYR	A	102	22.430	26.916	37.062	1.00	31.33	C
ATOM	697	CD2	TYR	A	102	23.842	28.844	37.054	1.00	31.29	C
ATOM	698	CE1	TYR	A	102	23.456	26.163	37.612	1.00	32.81	C
ATOM	699	CE2	TYR	A	102	24.869	28.098	37.601	1.00	32.24	C
ATOM	700	CZ	TYR	A	102	24.676	26.764	37.876	1.00	34.18	C
ATOM	701	OH	TYR	A	102	25.720	26.030	38.418	1.00	39.06	C
ATOM	702	N	TYR	A	103	21.684	29.478	39.488	1.00	31.28	O
ATOM	703	CA	TYR	A	103	22.569	29.983	40.539	1.00	31.56	N
ATOM	704	C	TYR	A	103	23.524	28.911	41.058	1.00	31.11	C
ATOM	705	O	TYR	A	103	23.190	27.732	41.165	1.00	30.75	C
ATOM	706	CB	TYR	A	103	21.796	30.599	41.706	1.00	31.98	O
ATOM	707	CG	TYR	A	103	20.846	29.663	42.385	1.00	33.92	C
ATOM	708	CD1	TYR	A	103	19.577	29.445	41.868	1.00	36.08	C
ATOM	709	CD2	TYR	A	103	21.203	29.010	43.553	1.00	36.91	C
ATOM	710	CE1	TYR	A	103	18.696	28.587	42.483	1.00	37.42	C
ATOM	711	CE2	TYR	A	103	20.325	28.153	44.189	1.00	37.63	C
ATOM	712	CZ	TYR	A	103	19.070	27.947	43.647	1.00	38.74	C
ATOM	713	OH	TYR	A	103	18.183	27.099	44.264	1.00	40.90	C
ATOM	714	N	ASP	A	104	24.725	29.345	41.391	1.00	30.34	O
ATOM	715	CA	ASP	A	104	25.752	28.444	41.873	1.00	30.04	N
ATOM	716	C	ASP	A	104	25.735	28.408	43.394	1.00	29.73	C
ATOM	717	O	ASP	A	104	26.079	29.388	44.062	1.00	28.53	C
ATOM	718	CB	ASP	A	104	27.089	28.931	41.350	1.00	30.07	O
ATOM	719	CG	ASP	A	104	28.233	28.037	41.731	1.00	30.02	C
ATOM	720	OD1	ASP	A	104	28.069	27.137	42.605	1.00	28.92	C
ATOM	721	OD2	ASP	A	104	29.339	28.183	41.168	1.00	29.24	O
ATOM	722	N	GLU	A	105	25.327	27.263	43.931	1.00	29.74	N

ATOM	723	CA	GLU	A	105	25.169	27.095	45.371	1.00	29.91
ATOM	724	C	GLU	A	105	26.461	27.343	46.155	1.00	29.29
ATOM	725	O	GLU	A	105	26.412	27.872	47.263	1.00	28.37
ATOM	726	CB	GLU	A	105	24.601	25.699	45.672	1.00	30.40
ATOM	727	CG	GLU	A	105	23.097	25.621	45.410	1.00	32.58
ATOM	728	CD	GLU	A	105	22.546	24.210	45.287	1.00	34.72
ATOM	729	OE1	GLU	A	105	22.945	23.320	46.072	1.00	35.38
ATOM	730	OE2	GLU	A	105	21.683	24.001	44.402	1.00	35.93
ATOM	731	N	LYS	A	106	27.607	27.000	45.565	1.00	29.02
ATOM	732	CA	LYS	A	106	28.897	27.156	46.243	1.00	29.26
ATOM	733	C	LYS	A	106	29.245	28.608	46.493	1.00	29.31
ATOM	734	O	LYS	A	106	30.051	28.903	47.363	1.00	28.83
ATOM	735	CB	LYS	A	106	30.036	26.530	45.433	1.00	29.60
ATOM	736	CG	LYS	A	106	29.864	25.036	45.094	1.00	30.18
ATOM	737	N	LYS	A	107	28.646	29.516	45.724	1.00	29.27
ATOM	738	CA	LYS	A	107	28.925	30.933	45.876	1.00	29.61
ATOM	739	C	LYS	A	107	27.933	31.605	46.820	1.00	30.29
ATOM	740	O	LYS	A	107	28.062	32.785	47.100	1.00	30.16
ATOM	741	CB	LYS	A	107	28.924	31.636	44.504	1.00	29.35
ATOM	742	CG	LYS	A	107	30.222	31.411	43.697	1.00	28.97
ATOM	743	CD	LYS	A	107	30.142	31.864	42.230	1.00	26.18
ATOM	744	CE	LYS	A	107	31.459	31.534	41.498	1.00	26.51
ATOM	745	NZ	LYS	A	107	31.578	32.052	40.083	1.00	24.21
ATOM	746	N	MET	A	108	26.950	30.869	47.320	1.00	31.39
ATOM	747	CA	MET	A	108	25.939	31.482	48.186	1.00	32.74
ATOM	748	C	MET	A	108	26.468	32.050	49.514	1.00	33.50
ATOM	749	O	MET	A	108	25.987	33.082	49.988	1.00	33.27
ATOM	750	CB	MET	A	108	24.787	30.509	48.417	1.00	33.11
ATOM	751	CG	MET	A	108	23.976	30.300	47.127	1.00	34.43
ATOM	752	SD	MET	A	108	22.458	29.330	47.263	1.00	37.47
ATOM	753	CE	MET	A	108	21.487	30.326	48.417	1.00	37.24
ATOM	754	N	ALA	A	109	27.484	31.427	50.090	1.00	34.73
ATOM	755	CA	ALA	A	109	28.039	31.922	51.359	1.00	36.13
ATOM	756	C	ALA	A	109	28.555	33.361	51.266	1.00	37.12
ATOM	757	O	ALA	A	109	28.455	34.127	52.213	1.00	37.35
ATOM	758	CB	ALA	A	109	29.147	30.991	51.860	1.00	35.96
ATOM	759	N	ASN	A	110	29.076	33.745	50.112	1.00	38.70
ATOM	760	CA	ASN	A	110	29.631	35.090	49.949	1.00	39.61
ATOM	761	C	ASN	A	110	28.605	36.188	49.621	1.00	39.61
ATOM	762	O	ASN	A	110	28.950	37.359	49.500	1.00	39.50
ATOM	763	CB	ASN	A	110	30.730	35.036	48.888	1.00	39.94
ATOM	764	CG	ASN	A	110	31.916	34.186	49.329	1.00	41.54
ATOM	765	OD1	ASN	A	110	32.341	34.223	50.503	1.00	41.36
ATOM	766	ND2	ASN	A	110	32.451	33.406	48.399	1.00	43.06
ATOM	767	N	PHE	A	111	27.349	35.802	49.465	1.00	40.12
ATOM	768	CA	PHE	A	111	26.278	36.753	49.208	1.00	40.43
ATOM	769	C	PHE	A	111	25.064	36.292	50.016	1.00	41.38
ATOM	770	O	PHE	A	111	24.068	35.811	49.471	1.00	40.89
ATOM	771	CB	PHE	A	111	25.954	36.839	47.710	1.00	40.30
ATOM	772	CG	PHE	A	111	26.973	37.597	46.910	1.00	38.60
ATOM	773	CD1	PHE	A	111	28.199	37.033	46.618	1.00	37.50
ATOM	774	CD2	PHE	A	111	26.705	38.867	46.445	1.00	38.02
ATOM	775	CE1	PHE	A	111	29.140	37.725	45.890	1.00	36.74
ATOM	776	CE2	PHE	A	111	27.649	39.560	45.705	1.00	36.60
ATOM	777	CZ	PHE	A	111	28.863	38.986	45.433	1.00	35.45
ATOM	778	N	GLN	A	112	25.175	36.457	51.329	1.00	42.72
ATOM	779	CA	GLN	A	112	24.154	36.022	52.276	1.00	43.86
ATOM	780	C	GLN	A	112	22.790	36.598	51.948	1.00	44.00
ATOM	781	O	GLN	A	112	21.774	35.994	52.280	1.00	44.39
ATOM	782	CB	GLN	A	112	24.535	36.428	53.706	1.00	44.29
ATOM	783	CG	GLN	A	112	25.923	35.985	54.166	1.00	46.45

C
C
C
C
C
C
O
O
N
C
C
O
C
C
N
C
C
N
N
C
C
C
O
C
C
S
C
N
C
C
C
C
O
C
C
C
O
N
N
C
C
C
O
C
C
C
C
C
C
N
C
C
C
O
C
C

ATOM	784	CD	GLN	A	112	26.050	34.477	54.296	1.00	49.14	C
ATOM	785	OE1	GLN	A	112	25.523	33.733	53.466	1.00	50.83	O
ATOM	786	NE2	GLN	A	112	26.756	34.022	55.332	1.00	50.52	N
ATOM	787	N	ASN	A	113	22.765	37.759	51.299	1.00	43.96	N
ATOM	788	CA	ASN	A	113	21.504	38.416	50.971	1.00	44.15	C
ATOM	789	C	ASN	A	113	20.827	37.921	49.679	1.00	44.12	C
ATOM	790	O	ASN	A	113	19.768	38.431	49.309	1.00	44.11	O
ATOM	791	CB	ASN	A	113	21.696	39.941	50.918	1.00	44.21	C
ATOM	792	CG	ASN	A	113	22.084	40.541	52.283	1.00	44.80	C
ATOM	793	OD1	ASN	A	113	21.759	39.991	53.349	1.00	43.65	O
ATOM	794	ND2	ASN	A	113	22.778	41.675	52.246	1.00	44.92	N
ATOM	795	N	PHE	A	114	21.415	36.939	48.995	1.00	43.98	N
ATOM	796	CA	PHE	A	114	20.793	36.411	47.787	1.00	43.71	C
ATOM	797	C	PHE	A	114	19.778	35.345	48.150	1.00	43.76	C
ATOM	798	O	PHE	A	114	20.111	34.362	48.815	1.00	43.32	O
ATOM	799	CB	PHE	A	114	21.813	35.808	46.833	1.00	43.82	C
ATOM	800	CG	PHE	A	114	21.184	35.128	45.650	1.00	43.80	C
ATOM	801	CD1	PHE	A	114	20.567	35.881	44.661	1.00	43.54	C
ATOM	802	CD2	PHE	A	114	21.170	33.745	45.541	1.00	43.65	C
ATOM	803	CE1	PHE	A	114	19.963	35.276	43.587	1.00	43.55	C
ATOM	804	CE2	PHE	A	114	20.566	33.129	44.451	1.00	43.21	C
ATOM	805	CZ	PHE	A	114	19.961	33.897	43.476	1.00	42.89	C
ATOM	806	N	LYS	A	115	18.543	35.543	47.705	1.00	43.96	N
ATOM	807	CA	LYS	A	115	17.459	34.613	47.999	1.00	44.38	C
ATOM	808	C	LYS	A	115	16.933	34.026	46.693	1.00	44.04	C
ATOM	809	O	LYS	A	115	16.213	34.687	45.945	1.00	43.97	O
ATOM	810	CB	LYS	A	115	16.342	35.329	48.761	1.00	44.79	C
ATOM	811	CG	LYS	A	115	16.763	35.866	50.129	1.00	46.74	C
ATOM	812	CD	LYS	A	115	17.021	34.744	51.130	1.00	48.83	C
ATOM	813	CE	LYS	A	115	17.461	35.288	52.484	1.00	50.18	C
ATOM	814	NZ	LYS	A	115	17.498	34.224	53.539	1.00	51.05	N
ATOM	815	N	PRO	A	116	17.270	32.771	46.433	1.00	43.65	N
ATOM	816	CA	PRO	A	116	16.918	32.133	45.160	1.00	43.43	C
ATOM	817	C	PRO	A	116	15.415	32.034	44.969	1.00	42.76	C
ATOM	818	O	PRO	A	116	14.711	31.724	45.915	1.00	42.78	O
ATOM	819	CB	PRO	A	116	17.525	30.732	45.273	1.00	43.57	C
ATOM	820	CG	PRO	A	116	18.385	30.743	46.496	1.00	43.88	C
ATOM	821	CD	PRO	A	116	17.948	31.855	47.359	1.00	43.68	C
ATOM	822	N	ARG	A	117	14.940	32.306	43.761	1.00	42.20	N
ATOM	823	CA	ARG	A	117	13.518	32.214	43.451	1.00	41.56	C
ATOM	824	C	ARG	A	117	13.140	30.780	43.104	1.00	41.00	C
ATOM	825	O	ARG	A	117	11.957	30.461	43.007	1.00	41.15	O
ATOM	826	CB	ARG	A	117	13.164	33.132	42.311	1.00	41.47	C
ATOM	827	N	SER	A	118	14.139	29.925	42.904	1.00	40.00	N
ATOM	828	CA	SER	A	118	13.882	28.517	42.654	1.00	39.60	C
ATOM	829	C	SER	A	118	14.700	27.655	43.621	1.00	39.30	C
ATOM	830	O	SER	A	118	15.756	28.078	44.087	1.00	39.24	O
ATOM	831	CB	SER	A	118	14.184	28.158	41.196	1.00	39.30	C
ATOM	832	OG	SER	A	118	15.560	28.228	40.942	1.00	38.53	O
ATOM	833	N	ASN	A	119	14.180	26.471	43.943	1.00	38.96	N
ATOM	834	CA	ASN	A	119	14.838	25.537	44.854	1.00	38.95	C
ATOM	835	C	ASN	A	119	15.115	24.215	44.160	1.00	37.99	C
ATOM	836	O	ASN	A	119	14.314	23.751	43.359	1.00	37.35	O
ATOM	837	CB	ASN	A	119	13.958	25.220	46.068	1.00	39.54	C
ATOM	838	CG	ASN	A	119	13.466	26.456	46.797	1.00	41.82	C
ATOM	839	OD1	ASN	A	119	14.255	27.279	47.289	1.00	44.50	O
ATOM	840	ND2	ASN	A	119	12.145	26.574	46.906	1.00	44.99	N
ATOM	841	N	ARG	A	120	16.243	23.607	44.492	1.00	37.46	N
ATOM	842	CA	ARG	A	120	16.627	22.325	43.927	1.00	37.15	C
ATOM	843	C	ARG	A	120	16.209	21.193	44.844	1.00	37.03	C
ATOM	844	O	ARG	A	120	16.359	21.272	46.069	1.00	37.28	O

ATOM	845	CB	ARG	A	120	18.144	22.264	43.730	1.00	37.07	C
ATOM	846	CG	ARG	A	120	18.632	21.020	42.993	1.00	36.32	C
ATOM	847	CD	ARG	A	120	20.152	20.886	42.938	1.00	35.46	C
ATOM	848	NE	ARG	A	120	20.777	21.277	44.198	1.00	34.29	N
ATOM	849	CZ	ARG	A	120	21.049	20.445	45.202	1.00	36.66	C
ATOM	850	NH1	ARG	A	120	20.753	19.147	45.122	1.00	35.60	N
ATOM	851	NH2	ARG	A	120	21.614	20.915	46.305	1.00	37.13	N
ATOM	852	N	GLU	A	121	15.665	20.141	44.250	1.00	36.76	N
ATOM	853	CA	GLU	A	121	15.326	18.944	44.993	1.00	36.29	C
ATOM	854	C	GLU	A	121	15.810	17.746	44.187	1.00	35.35	C
ATOM	855	O	GLU	A	121	15.709	17.726	42.953	1.00	35.38	O
ATOM	856	CB	GLU	A	121	13.820	18.863	45.262	1.00	36.78	C
ATOM	857	CG	GLU	A	121	13.398	17.585	45.971	1.00	39.29	C
ATOM	858	CD	GLU	A	121	12.126	17.732	46.795	1.00	42.09	C
ATOM	859	OE1	GLU	A	121	12.200	18.310	47.904	1.00	45.56	O
ATOM	860	OE2	GLU	A	121	11.059	17.254	46.350	1.00	43.04	O
ATOM	861	N	GLU	A	122	16.366	16.764	44.886	1.00	34.04	N
ATOM	862	CA	GLU	A	122	16.840	15.547	44.266	1.00	33.04	C
ATOM	863	C	GLU	A	122	15.748	14.518	44.415	1.00	32.55	C
ATOM	864	O	GLU	A	122	15.253	14.306	45.511	1.00	32.52	O
ATOM	865	CB	GLU	A	122	18.103	15.052	44.961	1.00	32.71	C
ATOM	866	CG	GLU	A	122	19.265	16.018	44.885	1.00	31.86	C
ATOM	867	CD	GLU	A	122	19.705	16.296	43.454	1.00	31.04	C
ATOM	868	OE1	GLU	A	122	20.065	15.331	42.757	1.00	29.68	O
ATOM	869	OE2	GLU	A	122	19.694	17.479	43.025	1.00	29.20	O
ATOM	870	N	MET	A	123	15.349	13.885	43.321	1.00	31.83	N
ATOM	871	CA	MET	A	123	14.329	12.858	43.415	1.00	31.40	C
ATOM	872	C	MET	A	123	14.532	11.820	42.343	1.00	30.68	C
ATOM	873	O	MET	A	123	15.380	11.982	41.457	1.00	30.07	O
ATOM	874	CB	MET	A	123	12.931	13.466	43.309	1.00	31.52	C
ATOM	875	CG	MET	A	123	12.667	14.205	42.032	1.00	32.93	C
ATOM	876	SD	MET	A	123	11.115	15.145	42.034	1.00	35.27	S
ATOM	877	CE	MET	A	123	11.554	16.565	42.966	1.00	35.33	C
ATOM	878	N	LYS	A	124	13.766	10.738	42.464	1.00	29.85	N
ATOM	879	CA	LYS	A	124	13.752	9.671	41.483	1.00	29.22	C
ATOM	880	C	LYS	A	124	12.891	10.140	40.307	1.00	28.45	C
ATOM	881	O	LYS	A	124	12.066	11.039	40.466	1.00	27.39	O
ATOM	882	CB	LYS	A	124	13.183	8.389	42.100	1.00	29.57	C
ATOM	883	CG	LYS	A	124	13.954	7.857	43.321	1.00	29.88	C
ATOM	884	CD	LYS	A	124	15.392	7.501	42.950	1.00	30.48	C
ATOM	885	CE	LYS	A	124	16.231	7.129	44.167	1.00	30.33	C
ATOM	886	NZ	LYS	A	124	17.691	7.276	43.872	1.00	30.47	N
ATOM	887	N	PHE	A	125	13.075	9.536	39.133	1.00	27.71	N
ATOM	888	CA	PHE	A	125	12.364	10.000	37.949	1.00	27.22	C
ATOM	889	C	PHE	A	125	10.869	9.891	38.114	1.00	27.29	C
ATOM	890	O	PHE	A	125	10.138	10.825	37.801	1.00	26.75	O
ATOM	891	CB	PHE	A	125	12.794	9.270	36.681	1.00	26.90	C
ATOM	892	CG	PHE	A	125	12.494	10.051	35.433	1.00	27.63	C
ATOM	893	CD1	PHE	A	125	13.330	11.083	35.032	1.00	27.70	C
ATOM	894	CD2	PHE	A	125	11.351	9.802	34.702	1.00	28.29	C
ATOM	895	CE1	PHE	A	125	13.059	11.828	33.905	1.00	27.96	C
ATOM	896	CE2	PHE	A	125	11.065	10.540	33.563	1.00	29.16	C
ATOM	897	CZ	PHE	A	125	11.924	11.568	33.167	1.00	28.71	C
ATOM	898	N	HIS	A	126	10.426	8.743	38.620	1.00	27.46	N
ATOM	899	CA	HIS	A	126	9.013	8.499	38.832	1.00	27.77	C
ATOM	900	C	HIS	A	126	8.430	9.522	39.794	1.00	27.71	C
ATOM	901	O	HIS	A	126	7.245	9.821	39.731	1.00	27.27	O
ATOM	902	CB	HIS	A	126	8.770	7.056	39.329	1.00	27.96	C
ATOM	903	CG	HIS	A	126	8.897	6.880	40.812	1.00	28.47	C
ATOM	904	ND1	HIS	A	126	7.831	7.026	41.672	1.00	29.29	N
ATOM	905	CD2	HIS	A	126	9.961	6.552	41.585	1.00	29.43	C

ATOM	906	CE1	HIS	A	126	8.236	6.812	42.912	1.00	30.26	C
ATOM	907	NE2	HIS	A	126	9.525	6.524	42.887	1.00	29.64	N
ATOM	908	N	GLU	A	127	9.262	10.059	40.679	1.00	28.09	N
ATOM	909	CA	GLU	A	127	8.803	11.078	41.615	1.00	28.68	C
ATOM	910	C	GLU	A	127	8.585	12.400	40.865	1.00	28.99	C
ATOM	911	O	GLU	A	127	7.626	13.129	41.123	1.00	29.15	O
ATOM	912	CB	GLU	A	127	9.795	11.233	42.775	1.00	28.52	C
ATOM	913	CG	GLU	A	127	9.931	9.968	43.626	1.00	29.84	C
ATOM	914	CD	GLU	A	127	10.873	10.121	44.810	1.00	30.26	C
ATOM	915	OE1	GLU	A	127	12.069	10.398	44.603	1.00	29.68	O
ATOM	916	OE2	GLU	A	127	10.410	9.943	45.962	1.00	32.95	O
ATOM	917	N	PHE	A	128	9.471	12.696	39.927	1.00	29.39	N
ATOM	918	CA	PHE	A	128	9.354	13.914	39.119	1.00	29.64	C
ATOM	919	C	PHE	A	128	8.047	13.859	38.333	1.00	30.31	C
ATOM	920	O	PHE	A	128	7.274	14.817	38.294	1.00	29.97	O
ATOM	921	CB	PHE	A	128	10.551	14.017	38.167	1.00	29.10	C
ATOM	922	CG	PHE	A	128	10.337	14.950	36.991	1.00	28.33	C
ATOM	923	CD1	PHE	A	128	10.106	16.301	37.188	1.00	26.53	C
ATOM	924	CD2	PHE	A	128	10.397	14.475	35.696	1.00	26.99	C
ATOM	925	CE1	PHE	A	128	9.920	17.149	36.120	1.00	27.50	C
ATOM	926	CE2	PHE	A	128	10.217	15.341	34.610	1.00	28.49	C
ATOM	927	CZ	PHE	A	128	9.976	16.668	34.823	1.00	26.78	C
ATOM	928	N	VAL	A	129	7.797	12.705	37.736	1.00	31.32	N
ATOM	929	CA	VAL	A	129	6.603	12.499	36.930	1.00	32.38	C
ATOM	930	C	VAL	A	129	5.338	12.658	37.775	1.00	32.96	C
ATOM	931	O	VAL	A	129	4.398	13.362	37.388	1.00	32.79	O
ATOM	932	CB	VAL	A	129	6.606	11.094	36.313	1.00	32.48	C
ATOM	933	CG1	VAL	A	129	5.313	10.840	35.551	1.00	32.96	C
ATOM	934	CG2	VAL	A	129	7.828	10.901	35.417	1.00	31.98	C
ATOM	935	N	GLU	A	130	5.329	11.993	38.925	1.00	33.61	N
ATOM	936	CA	GLU	A	130	4.205	12.067	39.853	1.00	34.29	C
ATOM	937	C	GLU	A	130	3.963	13.515	40.246	1.00	34.51	C
ATOM	938	O	GLU	A	130	2.832	13.986	40.220	1.00	33.80	O
ATOM	939	CB	GLU	A	130	4.481	11.206	41.087	1.00	34.35	C
ATOM	940	CG	GLU	A	130	4.372	9.711	40.815	1.00	34.92	C
ATOM	941	CD	GLU	A	130	5.204	8.858	41.761	1.00	35.66	C
ATOM	942	OE1	GLU	A	130	5.595	9.345	42.845	1.00	36.55	O
ATOM	943	OE2	GLU	A	130	5.477	7.692	41.407	1.00	36.06	O
ATOM	944	N	LYS	A	131	5.033	14.223	40.587	1.00	35.27	N
ATOM	945	CA	LYS	A	131	4.912	15.632	40.920	1.00	36.15	C
ATOM	946	C	LYS	A	131	4.286	16.408	39.758	1.00	36.63	C
ATOM	947	O	LYS	A	131	3.420	17.253	39.972	1.00	36.42	O
ATOM	948	CB	LYS	A	131	6.269	16.240	41.261	1.00	36.51	C
ATOM	949	CG	LYS	A	131	6.467	16.623	42.712	1.00	37.67	C
ATOM	950	CD	LYS	A	131	7.125	18.001	42.822	1.00	38.95	C
ATOM	951	CE	LYS	A	131	7.581	18.324	44.252	1.00	40.39	C
ATOM	952	NZ	LYS	A	131	8.073	19.747	44.404	1.00	40.82	N
ATOM	953	N	LEU	A	132	4.725	16.142	38.532	1.00	37.30	N
ATOM	954	CA	LEU	A	132	4.156	16.845	37.383	1.00	38.29	C
ATOM	955	C	LEU	A	132	2.669	16.557	37.280	1.00	38.83	C
ATOM	956	O	LEU	A	132	1.875	17.449	36.976	1.00	38.56	O
ATOM	957	CB	LEU	A	132	4.819	16.424	36.081	1.00	38.42	C
ATOM	958	CG	LEU	A	132	6.224	16.916	35.791	1.00	39.15	C
ATOM	959	CD1	LEU	A	132	6.671	16.296	34.485	1.00	39.76	C
ATOM	960	CD2	LEU	A	132	6.281	18.433	35.712	1.00	39.81	C
ATOM	961	N	GLN	A	133	2.311	15.300	37.529	1.00	39.68	N
ATOM	962	CA	GLN	A	133	0.920	14.863	37.495	1.00	40.45	C
ATOM	963	C	GLN	A	133	0.057	15.580	38.541	1.00	41.05	C
ATOM	964	O	GLN	A	133	-1.035	16.040	38.222	1.00	41.20	O
ATOM	965	CB	GLN	A	133	0.835	13.344	37.681	1.00	40.31	C
ATOM	966	N	ASP	A	134	0.542	15.682	39.777	1.00	41.89	N

ATOM	967	CA	ASP	A	134	-0.232	16.313	40.854	1.00	42.72	C
ATOM	968	C	ASP	A	134	-0.506	17.792	40.576	1.00	42.60	C
ATOM	969	O	ASP	A	134	-1.570	18.316	40.900	1.00	42.25	O
ATOM	970	CB	ASP	A	134	0.491	16.169	42.200	1.00	43.21	C
ATOM	971	CG	ASP	A	134	-0.429	16.419	43.396	1.00	45.71	C
ATOM	972	OD1	ASP	A	134	-1.566	16.916	43.213	1.00	48.39	O
ATOM	973	OD2	ASP	A	134	-0.104	16.131	44.571	1.00	49.60	O
ATOM	974	N	ILE	A	135	0.467	18.467	39.981	1.00	42.77	N
ATOM	975	CA	ILE	A	135	0.306	19.872	39.660	1.00	42.91	C
ATOM	976	C	ILE	A	135	-0.793	20.050	38.626	1.00	43.15	C
ATOM	977	O	ILE	A	135	-1.690	20.871	38.800	1.00	43.03	O
ATOM	978	CB	ILE	A	135	1.623	20.443	39.155	1.00	42.94	C
ATOM	979	CG1	ILE	A	135	2.586	20.607	40.330	1.00	43.10	C
ATOM	980	CG2	ILE	A	135	1.396	21.778	38.473	1.00	42.80	C
ATOM	981	CD1	ILE	A	135	4.040	20.621	39.934	1.00	43.49	C
ATOM	982	N	GLN	A	136	-0.716	19.272	37.554	1.00	43.64	N
ATOM	983	CA	GLN	A	136	-1.712	19.321	36.496	1.00	44.17	C
ATOM	984	C	GLN	A	136	-3.103	19.129	37.081	1.00	44.65	C
ATOM	985	O	GLN	A	136	-3.976	19.984	36.936	1.00	44.71	O
ATOM	986	CB	GLN	A	136	-1.431	18.240	35.466	1.00	44.19	C
ATOM	987	N	GLN	A	137	-3.289	18.014	37.776	1.00	45.23	N
ATOM	988	CA	GLN	A	137	-4.596	17.654	38.314	1.00	45.66	C
ATOM	989	C	GLN	A	137	-5.146	18.696	39.277	1.00	45.88	C
ATOM	990	O	GLN	A	137	-6.337	19.005	39.238	1.00	46.33	O
ATOM	991	CB	GLN	A	137	-4.535	16.277	38.991	1.00	45.74	C
ATOM	992	N	ARG	A	138	-4.288	19.246	40.131	1.00	45.93	N
ATOM	993	CA	ARG	A	138	-4.740	20.211	41.132	1.00	45.84	C
ATOM	994	C	ARG	A	138	-4.720	21.640	40.592	1.00	45.53	C
ATOM	995	O	ARG	A	138	-4.911	22.598	41.344	1.00	45.72	O
ATOM	996	CB	ARG	A	138	-3.880	20.108	42.398	1.00	45.92	C
ATOM	997	CG	ARG	A	138	-2.551	20.866	42.340	1.00	46.84	C
ATOM	998	CD	ARG	A	138	-1.589	20.458	43.437	1.00	47.70	C
ATOM	999	NE	ARG	A	138	-0.509	21.418	43.652	1.00	47.78	N
ATOM	1000	CZ	ARG	A	138	0.788	21.125	43.578	1.00	49.17	C
ATOM	1001	NH1	ARG	A	138	1.186	19.897	43.274	1.00	50.12	N
ATOM	1002	NH2	ARG	A	138	1.702	22.064	43.798	1.00	49.52	N
ATOM	1003	N	GLY	A	139	-4.492	21.783	39.290	1.00	44.99	N
ATOM	1004	CA	GLY	A	139	-4.419	23.094	38.669	1.00	44.52	C
ATOM	1005	C	GLY	A	139	-3.412	24.041	39.310	1.00	44.06	C
ATOM	1006	O	GLY	A	139	-3.551	25.259	39.203	1.00	44.28	O
ATOM	1007	N	GLY	A	140	-2.382	23.495	39.953	1.00	43.28	N
ATOM	1008	CA	GLY	A	140	-1.388	24.309	40.629	1.00	42.49	C
ATOM	1009	C	GLY	A	140	-0.609	25.218	39.694	1.00	41.92	C
ATOM	1010	O	GLY	A	140	-0.556	24.989	38.480	1.00	41.85	O
ATOM	1011	N	GLU	A	141	-0.004	26.264	40.250	1.00	40.90	N
ATOM	1012	CA	GLU	A	141	0.789	27.186	39.444	1.00	40.07	C
ATOM	1013	C	GLU	A	141	2.286	26.884	39.549	1.00	38.68	C
ATOM	1014	O	GLU	A	141	3.096	27.503	38.867	1.00	38.67	O
ATOM	1015	CB	GLU	A	141	0.513	28.644	39.839	1.00	40.36	C
ATOM	1016	CG	GLU	A	141	-0.799	29.224	39.309	1.00	41.91	C
ATOM	1017	CD	GLU	A	141	-1.001	29.042	37.805	1.00	44.25	C
ATOM	1018	OE1	GLU	A	141	0.004	29.008	37.043	1.00	45.69	O
ATOM	1019	OE2	GLU	A	141	-2.181	28.940	37.380	1.00	44.55	O
ATOM	1020	N	GLU	A	142	2.648	25.944	40.413	1.00	37.02	N
ATOM	1021	CA	GLU	A	142	4.040	25.573	40.596	1.00	35.61	C
ATOM	1022	C	GLU	A	142	4.629	25.120	39.265	1.00	34.41	C
ATOM	1023	O	GLU	A	142	3.923	24.594	38.408	1.00	34.25	O
ATOM	1024	CB	GLU	A	142	4.150	24.443	41.627	1.00	35.49	C
ATOM	1025	CG	GLU	A	142	5.571	24.173	42.095	1.00	35.47	C
ATOM	1026	CD	GLU	A	142	5.710	22.944	42.978	1.00	36.48	C
ATOM	1027	OE1	GLU	A	142	4.733	22.174	43.142	1.00	37.45	O

ATOM	1028	OE2	GLU	A	142	6.821	22.739	43.511	1.00	36.76
ATOM	1029	N	ARG	A	143	5.926	25.323	39.096	1.00	32.91
ATOM	1030	CA	ARG	A	143	6.596	24.893	37.884	1.00	31.95
ATOM	1031	C	ARG	A	143	7.803	24.058	38.214	1.00	30.62
ATOM	1032	O	ARG	A	143	8.514	24.354	39.166	1.00	30.75
ATOM	1033	CB	ARG	A	143	7.072	26.097	37.082	1.00	32.19
ATOM	1034	CG	ARG	A	143	5.968	26.912	36.447	1.00	32.57
ATOM	1035	CD	ARG	A	143	6.507	28.170	35.801	1.00	32.53
ATOM	1036	NE	ARG	A	143	5.492	28.885	35.037	1.00	32.25
ATOM	1037	CZ	ARG	A	143	5.158	28.617	33.790	1.00	31.80
ATOM	1038	NH1	ARG	A	143	5.757	27.637	33.105	1.00	31.28
ATOM	1039	NH2	ARG	A	143	4.214	29.345	33.221	1.00	32.77
ATOM	1040	N	LEU	A	144	8.054	23.028	37.417	1.00	28.97
ATOM	1041	CA	LEU	A	144	9.235	22.219	37.617	1.00	28.09
ATOM	1042	C	LEU	A	144	10.150	22.257	36.398	1.00	27.14
ATOM	1043	O	LEU	A	144	9.690	22.418	35.272	1.00	26.74
ATOM	1044	CB	LEU	A	144	8.834	20.783	37.887	1.00	28.28
ATOM	1045	CG	LEU	A	144	7.839	20.606	39.038	1.00	28.89
ATOM	1046	CD1	LEU	A	144	7.515	19.123	39.216	1.00	29.62
ATOM	1047	CD2	LEU	A	144	8.364	21.212	40.333	1.00	27.84
ATOM	1048	N	TYR	A	145	11.447	22.102	36.636	1.00	25.69
ATOM	1049	CA	TYR	A	145	12.395	21.976	35.548	1.00	25.05
ATOM	1050	C	TYR	A	145	13.459	20.957	35.968	1.00	25.00
ATOM	1051	O	TYR	A	145	14.239	21.193	36.881	1.00	25.01
ATOM	1052	CB	TYR	A	145	13.022	23.327	35.183	1.00	24.76
ATOM	1053	CG	TYR	A	145	13.471	23.485	33.728	1.00	23.25
ATOM	1054	CD1	TYR	A	145	13.739	22.385	32.931	1.00	21.70
ATOM	1055	CD2	TYR	A	145	13.667	24.746	33.175	1.00	22.16
ATOM	1056	CE1	TYR	A	145	14.154	22.526	31.620	1.00	20.98
ATOM	1057	CE2	TYR	A	145	14.097	24.904	31.846	1.00	21.30
ATOM	1058	CZ	TYR	A	145	14.332	23.799	31.071	1.00	20.66
ATOM	1059	OH	TYR	A	145	14.737	23.933	29.740	1.00	18.41
ATOM	1060	N	LEU	A	146	13.446	19.808	35.318	1.00	24.95
ATOM	1061	CA	LEU	A	146	14.453	18.787	35.547	1.00	25.43
ATOM	1062	C	LEU	A	146	15.678	19.100	34.694	1.00	24.91
ATOM	1063	O	LEU	A	146	15.555	19.313	33.493	1.00	24.10
ATOM	1064	CB	LEU	A	146	13.907	17.419	35.162	1.00	25.59
ATOM	1065	CG	LEU	A	146	14.875	16.238	35.334	1.00	27.52
ATOM	1066	CD1	LEU	A	146	14.111	14.966	35.672	1.00	26.95
ATOM	1067	CD2	LEU	A	146	15.742	16.023	34.084	1.00	29.46
ATOM	1068	N	GLN	A	147	16.845	19.097	35.330	1.00	24.91
ATOM	1069	CA	GLN	A	147	18.115	19.394	34.685	1.00	25.27
ATOM	1070	C	GLN	A	147	19.146	18.490	35.317	1.00	25.28
ATOM	1071	O	GLN	A	147	19.610	18.740	36.419	1.00	27.14
ATOM	1072	CB	GLN	A	147	18.475	20.876	34.863	1.00	25.02
ATOM	1073	CG	GLN	A	147	17.321	21.804	34.485	1.00	24.97
ATOM	1074	CD	GLN	A	147	17.682	23.283	34.560	1.00	27.08
ATOM	1075	OE1	GLN	A	147	16.891	24.149	34.141	1.00	27.93
ATOM	1076	NE2	GLN	A	147	18.860	23.575	35.064	1.00	23.50
ATOM	1077	N	GLN	A	148	19.487	17.428	34.611	1.00	25.47
ATOM	1078	CA	GLN	A	148	20.319	16.365	35.146	1.00	25.45
ATOM	1079	C	GLN	A	148	21.254	15.816	34.105	1.00	25.50
ATOM	1080	O	GLN	A	148	20.862	15.485	32.992	1.00	25.04
ATOM	1081	CB	GLN	A	148	19.436	15.231	35.640	1.00	25.37
ATOM	1082	CG	GLN	A	148	20.201	14.021	36.167	1.00	26.15
ATOM	1083	CD	GLN	A	148	21.129	14.383	37.314	1.00	26.56
ATOM	1084	OE1	GLN	A	148	20.718	15.093	38.240	1.00	23.94
ATOM	1085	NE2	GLN	A	148	22.383	13.916	37.248	1.00	25.29
ATOM	1086	N	THR	A	149	22.510	15.769	34.482	1.00	25.86
ATOM	1087	CA	THR	A	149	23.552	15.219	33.667	1.00	27.04
ATOM	1088	C	THR	A	149	23.298	13.720	33.527	1.00	27.25

O
N
C
C
O
C
C
C
N
C
N
N
N
C
C
O
C
C
C
C
C
C
C
C
O
N
C
C
C
O
C
C
C
C
O
N
N
C
C
C
O
N
N
C
C

ATOM	1089	O	THR	A	149	23.012	13.044	34.508	1.00	26.86	O
ATOM	1090	CB	THR	A	149	24.903	15.540	34.375	1.00	27.42	C
ATOM	1091	OG1	THR	A	149	25.300	16.882	34.020	1.00	29.80	O
ATOM	1092	CG2	THR	A	149	26.034	14.702	33.873	1.00	28.98	C
ATOM	1093	N	LEU	A	150	23.338	13.224	32.298	1.00	27.70	N
ATOM	1094	CA	LEU	A	150	23.195	11.812	32.027	1.00	28.46	C
ATOM	1095	C	LEU	A	150	24.429	11.076	32.573	1.00	28.97	C
ATOM	1096	O	LEU	A	150	25.548	11.468	32.268	1.00	29.00	O
ATOM	1097	CB	LEU	A	150	23.084	11.580	30.524	1.00	28.34	C
ATOM	1098	CG	LEU	A	150	21.780	11.981	29.837	1.00	29.63	C
ATOM	1099	CD1	LEU	A	150	21.944	11.862	28.328	1.00	30.03	C
ATOM	1100	CD2	LEU	A	150	20.619	11.140	30.310	1.00	31.15	C
ATOM	1101	N	ASN	A	151	24.230	10.030	33.378	1.00	29.35	N
ATOM	1102	CA	ASN	A	151	25.348	9.258	33.949	1.00	29.51	C
ATOM	1103	C	ASN	A	151	25.137	7.732	33.934	1.00	29.92	C
ATOM	1104	O	ASN	A	151	24.162	7.241	33.348	1.00	29.73	O
ATOM	1105	CB	ASN	A	151	25.590	9.710	35.385	1.00	29.78	C
ATOM	1106	CG	ASN	A	151	24.362	9.553	36.241	1.00	29.19	C
ATOM	1107	OD1	ASN	A	151	23.735	8.497	36.260	1.00	30.09	O
ATOM	1108	ND2	ASN	A	151	23.991	10.611	36.930	1.00	29.31	N
ATOM	1109	N	ASP	A	152	26.020	6.999	34.627	1.00	30.27	N
ATOM	1110	CA	ASP	A	152	26.034	5.514	34.649	1.00	30.85	C
ATOM	1111	C	ASP	A	152	24.830	4.797	35.212	1.00	30.51	C
ATOM	1112	O	ASP	A	152	24.779	3.562	35.137	1.00	30.29	O
ATOM	1113	CB	ASP	A	152	27.178	4.965	35.519	1.00	31.48	C
ATOM	1114	CG	ASP	A	152	28.384	5.812	35.493	1.00	34.64	C
ATOM	1115	OD1	ASP	A	152	28.588	6.523	34.482	1.00	41.71	O
ATOM	1116	OD2	ASP	A	152	29.177	5.857	36.437	1.00	37.31	O
ATOM	1117	N	THR	A	153	23.894	5.501	35.833	1.00	30.21	N
ATOM	1118	CA	THR	A	153	22.767	4.785	36.432	1.00	30.13	C
ATOM	1119	C	THR	A	153	21.612	4.626	35.463	1.00	29.54	C
ATOM	1120	O	THR	A	153	20.639	3.970	35.778	1.00	29.38	O
ATOM	1121	CB	THR	A	153	22.277	5.481	37.704	1.00	30.37	C
ATOM	1122	OG1	THR	A	153	21.735	6.770	37.373	1.00	31.06	O
ATOM	1123	CG2	THR	A	153	23.452	5.763	38.658	1.00	30.56	C
ATOM	1124	N	VAL	A	154	21.703	5.213	34.280	1.00	29.07	N
ATOM	1125	CA	VAL	A	154	20.596	5.082	33.346	1.00	28.90	C
ATOM	1126	C	VAL	A	154	20.488	3.613	32.968	1.00	28.83	C
ATOM	1127	O	VAL	A	154	21.486	2.898	33.002	1.00	28.78	O
ATOM	1128	CB	VAL	A	154	20.762	5.952	32.088	1.00	28.61	C
ATOM	1129	CG1	VAL	A	154	20.808	7.410	32.462	1.00	29.14	C
ATOM	1130	CG2	VAL	A	154	21.999	5.547	31.305	1.00	28.52	C
ATOM	1131	N	GLY	A	155	19.283	3.170	32.615	1.00	28.74	N
ATOM	1132	CA	GLY	A	155	19.043	1.780	32.263	1.00	28.33	C
ATOM	1133	C	GLY	A	155	19.484	1.387	30.860	1.00	28.70	C
ATOM	1134	O	GLY	A	155	19.862	2.235	30.031	1.00	28.27	O
ATOM	1135	N	ARG	A	156	19.384	0.088	30.591	1.00	28.47	N
ATOM	1136	CA	ARG	A	156	19.857	-0.519	29.349	1.00	28.54	C
ATOM	1137	C	ARG	A	156	19.291	0.096	28.084	1.00	28.18	C
ATOM	1138	O	ARG	A	156	20.029	0.346	27.143	1.00	28.69	O
ATOM	1139	CB	ARG	A	156	19.582	-2.019	29.362	1.00	28.58	C
ATOM	1140	N	LYS	A	157	17.986	0.304	28.042	1.00	27.53	N
ATOM	1141	CA	LYS	A	157	17.390	0.888	26.860	1.00	27.32	C
ATOM	1142	C	LYS	A	157	17.947	2.306	26.625	1.00	27.10	C
ATOM	1143	O	LYS	A	157	18.213	2.672	25.490	1.00	26.76	O
ATOM	1144	CB	LYS	A	157	15.858	0.902	26.960	1.00	27.68	C
ATOM	1145	CG	LYS	A	157	15.171	-0.451	26.656	1.00	26.26	C
ATOM	1146	N	ILE	A	158	18.138	3.086	27.688	1.00	26.68	N
ATOM	1147	CA	ILE	A	158	18.665	4.450	27.543	1.00	26.67	C
ATOM	1148	C	ILE	A	158	20.107	4.368	27.100	1.00	26.53	C
ATOM	1149	O	ILE	A	158	20.558	5.170	26.285	1.00	25.74	O

ATOM	1150	CB	ILE	A	158	18.570	5.245	28.837	1.00	26.51	C
ATOM	1151	CG1	ILE	A	158	17.114	5.399	29.269	1.00	27.11	C
ATOM	1152	CG2	ILE	A	158	19.208	6.618	28.667	1.00	27.55	C
ATOM	1153	CD1	ILE	A	158	16.232	6.028	28.260	1.00	28.98	C
ATOM	1154	N	VAL	A	159	20.817	3.378	27.629	1.00	26.46	N
ATOM	1155	CA	VAL	A	159	22.187	3.125	27.226	1.00	26.70	C
ATOM	1156	C	VAL	A	159	22.191	2.813	25.728	1.00	26.30	C
ATOM	1157	O	VAL	A	159	23.022	3.332	24.999	1.00	25.90	O
ATOM	1158	CB	VAL	A	159	22.819	1.963	28.018	1.00	27.18	C
ATOM	1159	CG1	VAL	A	159	24.045	1.436	27.311	1.00	28.12	C
ATOM	1160	CG2	VAL	A	159	23.180	2.410	29.427	1.00	27.44	C
ATOM	1161	N	MET	A	160	21.255	1.984	25.269	1.00	26.08	N
ATOM	1162	CA	MET	A	160	21.175	1.652	23.840	1.00	26.30	C
ATOM	1163	C	MET	A	160	20.906	2.929	23.005	1.00	24.71	C
ATOM	1164	O	MET	A	160	21.559	3.164	22.000	1.00	23.40	O
ATOM	1165	CB	MET	A	160	20.081	0.617	23.565	1.00	26.71	C
ATOM	1166	CG	MET	A	160	20.401	-0.790	24.070	1.00	30.23	C
ATOM	1167	SD	MET	A	160	21.721	-1.628	23.154	1.00	35.73	S
ATOM	1168	CE	MET	A	160	20.883	-1.823	21.524	1.00	36.68	C
ATOM	1169	N	ASP	A	161	19.948	3.739	23.439	1.00	23.86	N
ATOM	1170	CA	ASP	A	161	19.636	5.000	22.756	1.00	23.67	C
ATOM	1171	C	ASP	A	161	20.840	5.943	22.664	1.00	22.65	C
ATOM	1172	O	ASP	A	161	21.148	6.478	21.593	1.00	23.02	O
ATOM	1173	CB	ASP	A	161	18.472	5.699	23.455	1.00	23.42	C
ATOM	1174	CG	ASP	A	161	17.178	4.898	23.369	1.00	24.19	C
ATOM	1175	OD1	ASP	A	161	17.095	3.941	22.552	1.00	19.47	O
ATOM	1176	OD2	ASP	A	161	16.197	5.164	24.087	1.00	24.90	O
ATOM	1177	N	PHE	A	162	21.513	6.132	23.791	1.00	21.70	N
ATOM	1178	CA	PHE	A	162	22.667	7.003	23.903	1.00	21.16	C
ATOM	1179	C	PHE	A	162	23.777	6.557	22.964	1.00	20.02	C
ATOM	1180	O	PHE	A	162	24.393	7.362	22.310	1.00	20.22	O
ATOM	1181	CB	PHE	A	162	23.144	7.010	25.364	1.00	21.38	C
ATOM	1182	CG	PHE	A	162	24.286	7.936	25.650	1.00	22.82	C
ATOM	1183	CD1	PHE	A	162	24.072	9.290	25.882	1.00	26.05	C
ATOM	1184	CD2	PHE	A	162	25.579	7.450	25.738	1.00	24.99	C
ATOM	1185	CE1	PHE	A	162	25.144	10.136	26.181	1.00	25.32	C
ATOM	1186	CE2	PHE	A	162	26.654	8.301	26.026	1.00	24.73	C
ATOM	1187	CZ	PHE	A	162	26.438	9.622	26.250	1.00	24.82	C
ATOM	1188	N	LEU	A	163	24.029	5.271	22.894	1.00	19.74	N
ATOM	1189	CA	LEU	A	163	25.064	4.767	21.998	1.00	19.88	C
ATOM	1190	C	LEU	A	163	24.688	4.965	20.533	1.00	18.90	C
ATOM	1191	O	LEU	A	163	25.554	5.020	19.675	1.00	18.20	O
ATOM	1192	CB	LEU	A	163	25.320	3.287	22.256	1.00	19.78	C
ATOM	1193	CG	LEU	A	163	26.078	3.019	23.546	1.00	20.79	C
ATOM	1194	CD1	LEU	A	163	26.069	1.534	23.811	1.00	21.16	C
ATOM	1195	CD2	LEU	A	163	27.498	3.567	23.456	1.00	21.66	C
ATOM	1196	N	GLY	A	164	23.395	5.055	20.272	1.00	18.33	N
ATOM	1197	CA	GLY	A	164	22.883	5.296	18.941	1.00	19.05	C
ATOM	1198	C	GLY	A	164	22.879	6.764	18.519	1.00	19.33	C
ATOM	1199	O	GLY	A	164	22.347	7.079	17.451	1.00	20.16	O
ATOM	1200	N	PHE	A	165	23.424	7.651	19.350	1.00	18.45	N
ATOM	1201	CA	PHE	A	165	23.516	9.048	18.974	1.00	19.26	C
ATOM	1202	C	PHE	A	165	24.548	9.102	17.850	1.00	19.34	C
ATOM	1203	O	PHE	A	165	25.363	8.177	17.720	1.00	18.77	O
ATOM	1204	CB	PHE	A	165	23.947	9.915	20.158	1.00	19.09	C
ATOM	1205	CG	PHE	A	165	22.862	10.139	21.208	1.00	20.44	C
ATOM	1206	CD1	PHE	A	165	21.572	9.632	21.047	1.00	21.56	C
ATOM	1207	CD2	PHE	A	165	23.143	10.878	22.357	1.00	20.27	C
ATOM	1208	CE1	PHE	A	165	20.604	9.840	22.010	1.00	21.01	C
ATOM	1209	CE2	PHE	A	165	22.174	11.096	23.326	1.00	20.14	C
ATOM	1210	CZ	PHE	A	165	20.913	10.567	23.163	1.00	22.00	C

ATOM	1211	N	ASN	A	166	24.508	10.157	17.040	1.00	19.48
ATOM	1212	CA	ASN	A	166	25.428	10.290	15.897	1.00	20.08
ATOM	1213	C	ASN	A	166	26.827	10.792	16.305	1.00	20.53
ATOM	1214	O	ASN	A	166	27.193	11.975	16.076	1.00	20.34
ATOM	1215	CB	ASN	A	166	24.836	11.183	14.790	1.00	19.52
ATOM	1216	CG	ASN	A	166	25.572	11.003	13.451	1.00	19.82
ATOM	1217	OD1	ASN	A	166	26.652	10.376	13.412	1.00	18.39
ATOM	1218	ND2	ASN	A	166	24.987	11.528	12.350	1.00	15.25
ATOM	1219	N	TRP	A	167	27.564	9.892	16.955	1.00	20.69
ATOM	1220	CA	TRP	A	167	28.911	10.161	17.441	1.00	21.20
ATOM	1221	C	TRP	A	167	29.858	10.404	16.270	1.00	21.35
ATOM	1222	O	TRP	A	167	30.806	11.171	16.371	1.00	21.36
ATOM	1223	CB	TRP	A	167	29.402	8.977	18.316	1.00	21.31
ATOM	1224	CG	TRP	A	167	28.550	8.849	19.539	1.00	21.63
ATOM	1225	CD1	TRP	A	167	27.673	7.841	19.844	1.00	22.35
ATOM	1226	CD2	TRP	A	167	28.433	9.808	20.591	1.00	20.88
ATOM	1227	NE1	TRP	A	167	27.030	8.119	21.028	1.00	22.98
ATOM	1228	CE2	TRP	A	167	27.486	9.315	21.511	1.00	21.22
ATOM	1229	CE3	TRP	A	167	29.054	11.036	20.863	1.00	20.14
ATOM	1230	CZ2	TRP	A	167	27.143	9.999	22.670	1.00	22.17
ATOM	1231	CZ3	TRP	A	167	28.693	11.724	21.990	1.00	19.81
ATOM	1232	CH2	TRP	A	167	27.742	11.206	22.889	1.00	21.02
ATOM	1233	N	ASN	A	168	29.610	9.743	15.153	1.00	21.50
ATOM	1234	CA	ASN	A	168	30.464	9.927	13.991	1.00	21.77
ATOM	1235	C	ASN	A	168	30.488	11.406	13.605	1.00	21.48
ATOM	1236	O	ASN	A	168	31.549	11.992	13.428	1.00	20.55
ATOM	1237	CB	ASN	A	168	29.964	9.101	12.814	1.00	22.00
ATOM	1238	CG	ASN	A	168	30.856	9.236	11.584	1.00	23.99
ATOM	1239	OD1	ASN	A	168	32.052	8.974	11.661	1.00	26.79
ATOM	1240	ND2	ASN	A	168	30.277	9.652	10.448	1.00	24.40
ATOM	1241	N	TRP	A	169	29.302	11.995	13.484	1.00	21.28
ATOM	1242	CA	TRP	A	169	29.183	13.372	13.071	1.00	21.23
ATOM	1243	C	TRP	A	169	29.708	14.331	14.133	1.00	21.44
ATOM	1244	O	TRP	A	169	30.450	15.258	13.822	1.00	20.50
ATOM	1245	CB	TRP	A	169	27.736	13.741	12.740	1.00	21.29
ATOM	1246	CG	TRP	A	169	27.611	15.186	12.397	1.00	20.56
ATOM	1247	CD1	TRP	A	169	27.840	15.766	11.173	1.00	19.81
ATOM	1248	CD2	TRP	A	169	27.293	16.256	13.290	1.00	20.58
ATOM	1249	NE1	TRP	A	169	27.641	17.124	11.253	1.00	20.45
ATOM	1250	CE2	TRP	A	169	27.313	17.453	12.540	1.00	21.52
ATOM	1251	CE3	TRP	A	169	26.951	16.324	14.636	1.00	22.34
ATOM	1252	CZ2	TRP	A	169	27.036	18.705	13.102	1.00	24.24</

[illegible]

ATOM	1272	CA	LYS	A	172	34.255	14.963	13.592	1.00	26.46
ATOM	1273	C	LYS	A	172	34.221	16.478	13.873	1.00	25.77
ATOM	1274	O	LYS	A	172	35.224	17.169	13.681	1.00	25.85
ATOM	1275	CB	LYS	A	172	33.705	14.634	12.205	1.00	26.90
ATOM	1276	CG	LYS	A	172	34.262	13.351	11.668	1.00	29.88
ATOM	1277	CD	LYS	A	172	33.752	13.033	10.240	1.00	34.20
ATOM	1278	CE	LYS	A	172	34.458	11.810	9.690	1.00	36.15
ATOM	1279	NZ	LYS	A	172	34.556	11.800	8.188	1.00	40.56
ATOM	1280	N	GLN	A	173	33.088	16.986	14.338	1.00	25.37
ATOM	1281	CA	GLN	A	173	32.981	18.400	14.690	1.00	25.52
ATOM	1282	C	GLN	A	173	34.002	18.741	15.774	1.00	25.71
ATOM	1283	O	GLN	A	173	34.780	19.669	15.618	1.00	26.05
ATOM	1284	CB	GLN	A	173	31.589	18.740	15.200	1.00	25.30
ATOM	1285	CG	GLN	A	173	30.522	18.729	14.138	1.00	25.28
ATOM	1286	CD	GLN	A	173	30.783	19.753	13.061	1.00	26.34
ATOM	1287	OE1	GLN	A	173	31.252	20.872	13.345	1.00	26.57
ATOM	1288	NE2	GLN	A	173	30.492	19.386	11.822	1.00	23.68
ATOM	1289	N	GLN	A	174	33.972	17.985	16.867	1.00	25.20
ATOM	1290	CA	GLN	A	174	34.931	18.124	17.944	1.00	25.45
ATOM	1291	C	GLN	A	174	36.380	18.182	17.392	1.00	25.74
ATOM	1292	O	GLN	A	174	37.152	19.066	17.767	1.00	24.76
ATOM	1293	CB	GLN	A	174	34.770	16.951	18.923	1.00	25.13
ATOM	1294	CG	GLN	A	174	35.771	16.925	20.048	1.00	26.00
ATOM	1295	CD	GLN	A	174	35.636	15.697	20.929	1.00	26.76
ATOM	1296	OE1	GLN	A	174	35.278	14.629	20.450	1.00	28.02
ATOM	1297	NE2	GLN	A	174	35.917	15.851	22.222	1.00	26.75
ATOM	1298	N	GLY	A	175	36.724	17.249	16.505	1.00	25.61
ATOM	1299	CA	GLY	A	175	38.040	17.201	15.887	1.00	26.56
ATOM	1300	C	GLY	A	175	38.333	18.393	14.980	1.00	27.62
ATOM	1301	O	GLY	A	175	39.371	19.051	15.111	1.00	28.21
ATOM	1302	N	LYS	A	176	37.415	18.695	14.074	1.00	28.41
ATOM	1303	CA	LYS	A	176	37.583	19.834	13.173	1.00	29.52
ATOM	1304	C	LYS	A	176	37.778	21.177	13.895	1.00	29.38
ATOM	1305	O	LYS	A	176	38.557	22.009	13.443	1.00	29.30
ATOM	1306	CB	LYS	A	176	36.371	19.977	12.257	1.00	29.90
ATOM	1307	CG	LYS	A	176	36.212	18.895	11.227	1.00	32.69
ATOM	1308	CD	LYS	A	176	35.023	19.242	10.336	1.00	36.10
ATOM	1309	CE	LYS	A	176	34.114	18.065	10.148	1.00	37.89
ATOM	1310	NZ	LYS	A	176	32.736	18.469	9.746	1.00	40.12
ATOM	1311	N	ARG	A	177	37.060	21.397	14.993	1.00	29.00
ATOM	1312	CA	ARG	A	177	37.164	22.671	15.710	1.00	29.17
ATOM	1313	C	ARG	A	177	38.260	22.741	16.781	1.00	

C
C
O
C
C
C
C
N
N
C
C
O
C
C
C
C
O
N
N
C
C
O
C
C
C
C
O
N
N
C
C
O
C
C
C
C
C
C
C
C
N
N
C
C
O
C
C
C
C
C
C
N
N
C
C
O
C
C
C
C
C
N
N
C
C
O
C
C
C
C
C
C
C
C
C
C

ATOM	1333	CD2	TRP	A	179	33.985	22.140	19.825	1.00	24.35	
ATOM	1334	NE1	TRP	A	179	34.822	24.122	19.153	1.00	24.53	C
ATOM	1335	CE2	TRP	A	179	33.676	23.355	19.167	1.00	24.34	N
ATOM	1336	CE3	TRP	A	179	32.982	21.177	19.957	1.00	23.62	C
ATOM	1337	CZ2	TRP	A	179	32.417	23.627	18.659	1.00	22.63	C
ATOM	1338	CZ3	TRP	A	179	31.726	21.454	19.442	1.00	24.07	C
ATOM	1339	CH2	TRP	A	179	31.453	22.674	18.811	1.00	23.17	C
ATOM	1340	N	GLY	A	180	38.061	20.669	23.210	1.00	25.61	N
ATOM	1341	CA	GLY	A	180	38.471	19.737	24.240	1.00	25.44	C
ATOM	1342	C	GLY	A	180	37.428	18.634	24.377	1.00	26.09	C
ATOM	1343	O	GLY	A	180	36.592	18.432	23.491	1.00	26.60	O
ATOM	1344	N	GLN	A	181	37.448	17.934	25.503	1.00	26.33	N
ATOM	1345	CA	GLN	A	181	36.561	16.806	25.708	1.00	26.77	C
ATOM	1346	C	GLN	A	181	35.099	17.180	25.953	1.00	26.26	C
ATOM	1347	O	GLN	A	181	34.775	18.291	26.371	1.00	25.90	O
ATOM	1348	CB	GLN	A	181	37.044	15.970	26.887	1.00	27.12	C
ATOM	1349	CG	GLN	A	181	36.689	16.543	28.269	1.00	29.66	C
ATOM	1350	CD	GLN	A	181	36.917	15.533	29.388	1.00	33.17	C
ATOM	1351	OE1	GLN	A	181	38.055	15.295	29.791	1.00	35.52	O
ATOM	1352	NE2	GLN	A	181	35.839	14.920	29.869	1.00	35.59	N
ATOM	1353	N	LEU	A	182	34.238	16.225	25.646	1.00	25.83	N
ATOM	1354	CA	LEU	A	182	32.831	16.241	26.021	1.00	25.59	C
ATOM	1355	C	LEU	A	182	32.850	16.162	27.543	1.00	24.45	C
ATOM	1356	O	LEU	A	182	33.433	15.209	28.084	1.00	23.27	O
ATOM	1357	CB	LEU	A	182	32.180	14.958	25.499	1.00	25.74	C
ATOM	1358	CG	LEU	A	182	30.666	14.790	25.403	1.00	28.29	C
ATOM	1359	CD1	LEU	A	182	30.276	13.328	25.676	1.00	27.13	C
ATOM	1360	CD2	LEU	A	182	29.928	15.671	26.303	1.00	30.64	C
ATOM	1361	N	THR	A	183	32.288	17.158	28.239	1.00	23.24	N
ATOM	1362	CA	THR	A	183	32.256	17.110	29.699	1.00	22.37	C
ATOM	1363	C	THR	A	183	30.956	16.504	30.150	1.00	22.01	C
ATOM	1364	O	THR	A	183	30.907	15.867	31.186	1.00	21.99	O
ATOM	1365	CB	THR	A	183	32.395	18.504	30.371	1.00	22.62	C
ATOM	1366	OG1	THR	A	183	31.367	19.393	29.893	1.00	20.93	O
ATOM	1367	CG2	THR	A	183	33.707	19.154	30.013	1.00	22.33	C
ATOM	1368	N	SER	A	184	29.885	16.711	29.396	1.00	21.54	N
ATOM	1369	CA	SER	A	184	28.622	16.146	29.810	1.00	21.58	C
ATOM	1370	C	SER	A	184	27.498	16.381	28.866	1.00	21.50	C
ATOM	1371	O	SER	A	184	27.610	17.168	27.928	1.00	21.85	O
ATOM	1372	CB	SER	A	184	28.209	16.715	31.158	1.00	21.78	C
ATOM	1373	OG	SER	A	184	27.856	18.072	31.056	1.00	23.20	O
ATOM	1374	N	ASN	A	185	26.416	15.667	29.134	1.00	21.14	N
ATOM	1375	CA	ASN	A	185	25.167	15.807	28.427	1.00	22.40	C
ATOM	1376	C	ASN	A	185	24.104	16.059	29.459	1.00	22.16	C
ATOM	1377	O	ASN	A	185	23.802	15.189	30.272	1.00	22.11	O
ATOM	1378	CB	ASN	A	185	24.794	14.523	27.664	1.00	22.46	C
ATOM	1379	CG	ASN	A	185	25.798	14.174	26.604	1.00	23.64	C
ATOM	1380	OD1	ASN	A	185	26.582	13.257	26.789	1.00	25.26	O
ATOM	1381	ND2	ASN	A	185	25.794	14.907	25.494	1.00	22.02	N
ATOM	1382	N	LEU	A	186	23.507	17.228	29.391	1.00	22.02	N
ATOM	1383	CA	LEU	A	186	22.461	17.600	30.319	1.00	22.34	C
ATOM	1384	C	LEU	A	186	21.114	17.251	29.707	1.00	22.21	C
ATOM	1385	O	LEU	A	186	20.808	17.623	28.582	1.00	22.59	O
ATOM	1386	CB	LEU	A	186	22.519	19.111	30.591	1.00	22.36	C
ATOM	1387	CG	LEU	A	186	21.650	19.642	31.739	1.00	23.03	C
ATOM	1388	CD1	LEU	A	186	22.124	19.073	33.069	1.00	22.66	C
ATOM	1389	CD2	LEU	A	186	21.636	21.213	31.782	1.00	21.88	C
ATOM	1390	N	LEU	A	187	20.307	16.530	30.461	1.00	22.39	N
ATOM	1391	CA	LEU	A	187	18.949	16.234	30.057	1.00	21.93	C
ATOM	1392	C	LEU	A	187	18.089	17.336	30.659	1.00	21.94	C
ATOM	1393	O	LEU	A	187	18.163	17.569	31.870	1.00	21.56	O

ATOM	1394	CB	LEU	A	187	18.525	14.883	30.604	1.00	21.36
ATOM	1395	CG	LEU	A	187	17.037	14.555	30.489	1.00	22.46
ATOM	1396	CD1	LEU	A	187	16.550	14.530	29.038	1.00	22.25
ATOM	1397	CD2	LEU	A	187	16.783	13.200	31.154	1.00	22.25
ATOM	1398	N	LEU	A	188	17.301	18.028	29.826	1.00	21.68
ATOM	1399	CA	LEU	A	188	16.400	19.052	30.329	1.00	22.46
ATOM	1400	C	LEU	A	188	14.925	18.743	30.021	1.00	22.70
ATOM	1401	O	LEU	A	188	14.511	18.622	28.864	1.00	23.01
ATOM	1402	CB	LEU	A	188	16.747	20.432	29.769	1.00	22.51
ATOM	1403	CG	LEU	A	188	18.166	20.932	29.998	1.00	24.09
ATOM	1404	CD1	LEU	A	188	18.916	20.996	28.697	1.00	26.99
ATOM	1405	CD2	LEU	A	188	18.135	22.308	30.564	1.00	25.94
ATOM	1406	N	ILE	A	189	14.117	18.652	31.061	1.00	22.31
ATOM	1407	CA	ILE	A	189	12.721	18.383	30.851	1.00	22.23
ATOM	1408	C	ILE	A	189	11.959	19.459	31.554	1.00	22.43
ATOM	1409	O	ILE	A	189	12.045	19.599	32.773	1.00	22.44
ATOM	1410	CB	ILE	A	189	12.328	17.009	31.369	1.00	21.56
ATOM	1411	CG1	ILE	A	189	13.178	15.944	30.695	1.00	21.26
ATOM	1412	CG2	ILE	A	189	10.873	16.794	31.078	1.00	21.74
ATOM	1413	CD1	ILE	A	189	12.851	14.481	31.136	1.00	22.52
ATOM	1414	N	GLY	A	190	11.226	20.239	30.770	1.00	23.11
ATOM	1415	CA	GLY	A	190	10.545	21.396	31.294	1.00	23.06
ATOM	1416	C	GLY	A	190	9.084	21.395	30.988	1.00	23.56
ATOM	1417	O	GLY	A	190	8.594	20.646	30.117	1.00	23.29
ATOM	1418	N	MET	A	191	8.385	22.224	31.755	1.00	24.04
ATOM	1419	CA	MET	A	191	6.980	22.468	31.542	1.00	24.73
ATOM	1420	C	MET	A	191	6.837	23.623	30.576	1.00	24.58
ATOM	1421	O	MET	A	191	7.747	24.464	30.443	1.00	24.72
ATOM	1422	CB	MET	A	191	6.301	22.821	32.860	1.00	25.52
ATOM	1423	CG	MET	A	191	6.212	21.649	33.806	1.00	26.99
ATOM	1424	SD	MET	A	191	5.710	22.122	35.447	1.00	30.62
ATOM	1425	CE	MET	A	191	4.076	22.639	35.135	1.00	32.05
ATOM	1426	N	GLU	A	192	5.701	23.659	29.893	1.00	24.37
ATOM	1427	CA	GLU	A	192	5.407	24.718	28.940	1.00	24.51
ATOM	1428	C	GLU	A	192	5.468	26.072	29.628	1.00	24.20
ATOM	1429	O	GLU	A	192	5.013	26.237	30.745	1.00	24.12
ATOM	1430	CB	GLU	A	192	4.029	24.514	28.342	1.00	24.59
ATOM	1431	CG	GLU	A	192	2.930	24.487	29.384	1.00	26.08
ATOM	1432	CD	GLU	A	192	1.590	24.051	28.830	1.00	26.38
ATOM	1433	OE1	GLU	A	192	1.530	23.568	27.676	1.00	27.04
ATOM	1434	OE2	GLU	A	192	0.599	24.219	29.562	1.00	25.46
ATOM	1435	N	GLY	A	193	6.045	27.051	28.962	1.00	24.44
ATOM	1436	CA	GLY	A	193	6.153	28.364	29.562	1.00	24.21
ATOM	1437	C	GLY	A	193	7.428	28.565	30.358	1.00	23.71
ATOM	1438	O	GLY	A	193	7.728	29.691	30.697	1.00	24.97
ATOM	1439	N	ASN	A	194	8.181	27.507	30.663	1.00	23.27
ATOM	1440	CA	ASN	A	194	9.445	27.656	31.413	1.00	22.47
ATOM	1441	C	ASN	A	194	10.465	28.476	30.636	1.00	22.35
ATOM	1442	O	ASN	A	194	10.594	28.317	29.416	1.00	22.38
ATOM	1443	CB	ASN	A	194	10.108	26.299	31.708	1.00	22.17
ATOM	1444	CG	ASN	A	194	9.476	25.555	32.882	1.00	22.24
ATOM	1445	OD1	ASN	A	194	8.477	25.992	33.453	1.00	23.81
ATOM	1446	ND2	ASN	A	194	10.075	24.429	33.253	1.00	18.58
ATOM	1447	N	VAL	A	195	11.226	29.295	31.362	1.00	21.64
ATOM	1448	CA	VAL	A	195	12.263	30.114	30.789	1.00	21.41
ATOM	1449	C	VAL	A	195	13.569	29.988	31.556	1.00	20.40
ATOM	1450	O	VAL	A	195	13.603	30.014	32.790	1.00	20.03
ATOM	1451	CB	VAL	A	195	11.889	31.621	30.828	1.00	21.72
ATOM	1452	CG1	VAL	A	195	13.038	32.476	30.327	1.00	21.84
ATOM	1453	CG2	VAL	A	195	10.628	31.904	30.043	1.00	23.61
ATOM	1454	N	THR	A	196	14.644	29.834	30.813	1.00	19.77

C
C
C
C
N
C
C
O
C
C
C
C
N
C
C
O
C
C
N
C
C
O
N
C
C
C
O
C
C
S
C
N
C
C
C
O
C
C
O
O
N
C
C
O
N
C
C
O
C
C
O
N
N
C
C
C
C
N

ATOM	1455	CA	THR	A	196	15.980	29.917	31.376	1.00	20.37
ATOM	1456	C	THR	A	196	16.467	31.291	30.933	1.00	20.92
ATOM	1457	O	THR	A	196	16.649	31.505	29.724	1.00	20.45
ATOM	1458	CB	THR	A	196	16.882	28.865	30.793	1.00	19.87
ATOM	1459	OG1	THR	A	196	16.457	27.564	31.227	1.00	21.07
ATOM	1460	CG2	THR	A	196	18.273	29.043	31.340	1.00	20.00
ATOM	1461	N	PRO	A	197	16.602	32.225	31.879	1.00	21.53
ATOM	1462	CA	PRO	A	197	16.997	33.609	31.572	1.00	22.24
ATOM	1463	C	PRO	A	197	18.375	33.688	30.952	1.00	22.26
ATOM	1464	O	PRO	A	197	19.195	32.773	31.130	1.00	22.66
ATOM	1465	CB	PRO	A	197	16.998	34.305	32.934	1.00	22.51
ATOM	1466	CG	PRO	A	197	16.282	33.411	33.839	1.00	22.92
ATOM	1467	CD	PRO	A	197	16.367	32.028	33.312	1.00	21.85
ATOM	1468	N	ALA	A	198	18.606	34.776	30.234	1.00	21.54
ATOM	1469	CA	ALA	A	198	19.821	34.976	29.486	1.00	21.47
ATOM	1470	C	ALA	A	198	21.110	34.764	30.282	1.00	21.22
ATOM	1471	O	ALA	A	198	21.298	35.310	31.386	1.00	20.56
ATOM	1472	CB	ALA	A	198	19.809	36.397	28.891	1.00	21.92
ATOM	1473	N	HIS	A	199	22.023	34.020	29.683	1.00	21.01
ATOM	1474	CA	HIS	A	199	23.321	33.764	30.288	1.00	21.40
ATOM	1475	C	HIS	A	199	24.250	33.279	29.213	1.00	21.61
ATOM	1476	O	HIS	A	199	23.811	33.017	28.089	1.00	21.18
ATOM	1477	CB	HIS	A	199	23.217	32.669	31.338	1.00	21.05
ATOM	1478	CG	HIS	A	199	22.864	31.354	30.746	1.00	22.98
ATOM	1479	ND1	HIS	A	199	21.576	31.048	30.370	1.00	21.97
ATOM	1480	CD2	HIS	A	199	23.630	30.287	30.399	1.00	21.48
ATOM	1481	CE1	HIS	A	199	21.558	29.838	29.838	1.00	22.85
ATOM	1482	NE2	HIS	A	199	22.791	29.368	29.821	1.00	24.36
ATOM	1483	N	TYR	A	200	25.538	33.177	29.555	1.00	22.22
ATOM	1484	CA	TYR	A	200	26.531	32.597	28.654	1.00	22.56
ATOM	1485	C	TYR	A	200	27.234	31.448	29.380	1.00	22.96
ATOM	1486	O	TYR	A	200	27.293	31.430	30.612	1.00	22.90
ATOM	1487	CB	TYR	A	200	27.528	33.610	28.129	1.00	22.56
ATOM	1488	CG	TYR	A	200	28.492	34.228	29.145	1.00	23.02
ATOM	1489	CD1	TYR	A	200	29.731	33.671	29.385	1.00	22.79
ATOM	1490	CD2	TYR	A	200	28.180	35.419	29.799	1.00	23.89
ATOM	1491	CE1	TYR	A	200	30.612	34.229	30.283	1.00	22.32
ATOM	1492	CE2	TYR	A	200	29.062	35.991	30.722	1.00	22.06
ATOM	1493	CZ	TYR	A	200	30.277	35.402	30.947	1.00	22.13
ATOM	1494	OH	TYR	A	200	31.163	35.957	31.843	1.00	20.46
ATOM	1495	N	ASP	A	201	27.723	30.467	28.619	1.00	23.11
ATOM	1496	CA	ASP	A	201	28.433	29.326	29.213	1.00	23.02
ATOM	1497	C	ASP	A	201	29.833	2			

ATOM	1516	CB	GLN	A	203	31.373	24.363	27.618	1.00	23.43	C
ATOM	1517	CG	GLN	A	203	32.256	24.031	28.826	1.00	23.66	C
ATOM	1518	CD	GLN	A	203	32.061	22.612	29.354	1.00	25.58	C
ATOM	1519	OE1	GLN	A	203	31.915	21.661	28.582	1.00	27.85	O
ATOM	1520	NE2	GLN	A	203	32.084	22.468	30.670	1.00	25.67	N
ATOM	1521	N	GLN	A	204	31.465	25.013	24.575	1.00	22.51	N
ATOM	1522	CA	GLN	A	204	30.791	24.910	23.299	1.00	22.51	C
ATOM	1523	C	GLN	A	204	29.626	23.980	23.504	1.00	21.70	C
ATOM	1524	O	GLN	A	204	29.737	23.026	24.255	1.00	21.24	O
ATOM	1525	CB	GLN	A	204	31.718	24.307	22.255	1.00	22.29	C
ATOM	1526	CG	GLN	A	204	33.001	25.081	22.072	1.00	22.69	C
ATOM	1527	CD	GLN	A	204	32.820	26.346	21.259	1.00	22.39	C
ATOM	1528	OE1	GLN	A	204	31.699	26.736	20.932	1.00	19.22	O
ATOM	1529	NE2	GLN	A	204	33.937	26.970	20.906	1.00	19.77	N
ATOM	1530	N	ASN	A	205	28.523	24.241	22.810	1.00	21.48	N
ATOM	1531	CA	ASN	A	205	27.309	23.471	23.017	1.00	21.04	C
ATOM	1532	C	ASN	A	205	26.558	23.136	21.730	1.00	21.10	C
ATOM	1533	O	ASN	A	205	26.188	24.043	20.949	1.00	20.02	O
ATOM	1534	CB	ASN	A	205	26.401	24.322	23.920	1.00	21.70	C
ATOM	1535	CG	ASN	A	205	25.084	23.654	24.269	1.00	21.87	C
ATOM	1536	OD1	ASN	A	205	24.732	22.577	23.775	1.00	22.07	O
ATOM	1537	ND2	ASN	A	205	24.332	24.320	25.118	1.00	19.37	N
ATOM	1538	N	PHE	A	206	26.350	21.835	21.497	1.00	20.28	N
ATOM	1539	CA	PHE	A	206	25.372	21.414	20.523	1.00	20.14	C
ATOM	1540	C	PHE	A	206	24.141	21.030	21.323	1.00	19.94	C
ATOM	1541	O	PHE	A	206	24.164	20.052	22.065	1.00	20.71	O
ATOM	1542	CB	PHE	A	206	25.851	20.238	19.693	1.00	20.65	C
ATOM	1543	CG	PHE	A	206	26.799	20.618	18.620	1.00	20.30	C
ATOM	1544	CD1	PHE	A	206	26.394	21.438	17.600	1.00	20.89	C
ATOM	1545	CD2	PHE	A	206	28.101	20.168	18.642	1.00	21.26	C
ATOM	1546	CE1	PHE	A	206	27.254	21.783	16.605	1.00	22.17	C
ATOM	1547	CE2	PHE	A	206	28.980	20.523	17.641	1.00	21.31	C
ATOM	1548	CZ	PHE	A	206	28.558	21.336	16.632	1.00	21.79	C
ATOM	1549	N	PHE	A	207	23.058	21.764	21.081	1.00	19.87	N
ATOM	1550	CA	PHE	A	207	21.790	21.723	21.824	1.00	20.08	C
ATOM	1551	C	PHE	A	207	20.766	20.988	20.975	1.00	20.29	C
ATOM	1552	O	PHE	A	207	20.276	21.519	20.011	1.00	19.94	O
ATOM	1553	CB	PHE	A	207	21.385	23.190	22.083	1.00	20.46	C
ATOM	1554	CG	PHE	A	207	20.129	23.404	22.890	1.00	19.14	C
ATOM	1555	CD1	PHE	A	207	18.947	23.708	22.267	1.00	19.77	C
ATOM	1556	CD2	PHE	A	207	20.169	23.428	24.259	1.00	19.99	C
ATOM	1557	CE1	PHE	A	207	17.818	23.980	22.984	1.00	21.67	C
ATOM	1558	CE2	PHE	A	207	19.037	23.683	24.992	1.00	22.36	C
ATOM	1559	CZ	PHE	A	207	17.854	23.975	24.351	1.00	22.30	C
ATOM	1560	N	ALA	A	208	20.480	19.746	21.352	1.00	21.01	N
ATOM	1561	CA	ALA	A	208	19.688	18.827	20.541	1.00	21.26	C
ATOM	1562	C	ALA	A	208	18.260	18.669	21.030	1.00	21.74	C
ATOM	1563	O	ALA	A	208	18.000	17.983	22.049	1.00	21.21	O
ATOM	1564	CB	ALA	A	208	20.365	17.464	20.557	1.00	20.94	C
ATOM	1565	N	GLN	A	209	17.329	19.245	20.276	1.00	21.56	N
ATOM	1566	CA	GLN	A	209	15.940	19.219	20.697	1.00	21.78	C
ATOM	1567	C	GLN	A	209	15.289	17.864	20.393	1.00	21.84	C
ATOM	1568	O	GLN	A	209	15.506	17.245	19.323	1.00	20.40	O
ATOM	1569	CB	GLN	A	209	15.203	20.391	20.068	1.00	22.08	C
ATOM	1570	CG	GLN	A	209	13.790	20.597	20.542	1.00	22.02	C
ATOM	1571	CD	GLN	A	209	13.688	20.986	22.013	1.00	22.84	C
ATOM	1572	OE1	GLN	A	209	14.700	21.240	22.678	1.00	22.55	O
ATOM	1573	NE2	GLN	A	209	12.448	21.034	22.524	1.00	21.97	N
ATOM	1574	N	ILE	A	210	14.480	17.418	21.351	1.00	22.17	N
ATOM	1575	CA	ILE	A	210	13.904	16.084	21.329	1.00	22.53	C
ATOM	1576	C	ILE	A	210	12.397	16.074	21.316	1.00	23.45	C

ATOM	1577	O	ILE	A	210	11.813	15.429	20.467	1.00	24.08
ATOM	1578	CB	ILE	A	210	14.411	15.306	22.541	1.00	22.81
ATOM	1579	CG1	ILE	A	210	15.857	14.886	22.284	1.00	23.77
ATOM	1580	CG2	ILE	A	210	13.573	14.064	22.785	1.00	23.17
ATOM	1581	CD1	ILE	A	210	16.631	14.512	23.519	1.00	24.86
ATOM	1582	N	LYS	A	211	11.772	16.778	22.258	1.00	23.73
ATOM	1583	CA	LYS	A	211	10.318	16.829	22.351	1.00	22.99
ATOM	1584	C	LYS	A	211	9.873	18.253	22.566	1.00	22.73
ATOM	1585	O	LYS	A	211	10.436	18.969	23.391	1.00	22.34
ATOM	1586	CB	LYS	A	211	9.818	16.014	23.527	1.00	23.40
ATOM	1587	CG	LYS	A	211	8.285	15.757	23.509	1.00	24.55
ATOM	1588	CD	LYS	A	211	7.794	15.232	24.870	1.00	25.75
ATOM	1589	CE	LYS	A	211	6.389	14.579	24.861	1.00	25.61
ATOM	1590	NZ	LYS	A	211	5.580	14.641	23.613	1.00	25.29
ATOM	1591	N	GLY	A	212	8.832	18.655	21.847	1.00	22.34
ATOM	1592	CA	GLY	A	212	8.292	19.989	21.983	1.00	22.46
ATOM	1593	C	GLY	A	212	9.188	21.026	21.338	1.00	22.40
ATOM	1594	O	GLY	A	212	10.172	20.705	20.650	1.00	21.93
ATOM	1595	N	TYR	A	213	8.859	22.281	21.592	1.00	22.95
ATOM	1596	CA	TYR	A	213	9.530	23.400	20.944	1.00	23.32
ATOM	1597	C	TYR	A	213	10.079	24.412	21.926	1.00	23.27
ATOM	1598	O	TYR	A	213	9.434	24.740	22.918	1.00	23.34
ATOM	1599	CB	TYR	A	213	8.547	24.092	20.012	1.00	23.77
ATOM	1600	CG	TYR	A	213	8.133	23.198	18.887	1.00	25.58
ATOM	1601	CD1	TYR	A	213	7.110	22.261	19.041	1.00	28.74
ATOM	1602	CD2	TYR	A	213	8.798	23.253	17.687	1.00	28.55
ATOM	1603	CE1	TYR	A	213	6.764	21.410	17.995	1.00	30.77
ATOM	1604	CE2	TYR	A	213	8.465	22.429	16.651	1.00	30.41
ATOM	1605	CZ	TYR	A	213	7.461	21.516	16.795	1.00	31.44
ATOM	1606	OH	TYR	A	213	7.188	20.720	15.708	1.00	35.50
ATOM	1607	N	LYS	A	214	11.270	24.911	21.620	1.00	22.60
ATOM	1608	CA	LYS	A	214	11.916	25.911	22.431	1.00	22.58
ATOM	1609	C	LYS	A	214	12.353	27.077	21.576	1.00	22.52
ATOM	1610	O	LYS	A	214	12.937	26.898	20.493	1.00	23.74
ATOM	1611	CB	LYS	A	214	13.132	25.345	23.169	1.00	22.31
ATOM	1612	CG	LYS	A	214	12.784	24.539	24.392	1.00	22.79
ATOM	1613	CD	LYS	A	214	14.053	24.007	25.077	1.00	22.96
ATOM	1614	CE	LYS	A	214	13.821	23.675	26.530	1.00	20.69
ATOM	1615	NZ	LYS	A	214	15.076	23.085	27.149	1.00	19.14
ATOM	1616	N	ARG	A	215	12.037	28.277	22.047	1.00	21.95
ATOM	1617	CA	ARG	A	215	12.480	29.482	21.378	1.00	21.48
ATOM	1618	C	ARG	A	215	13.790	29.864	22.018	1.00	21.06
ATOM	1619	O	ARG	A	215	13.890	29.954	23.230	1.00	20.53
ATOM	1620	CB	ARG	A	215	11.466	30.582	21.576	1.00	22.15
ATOM	1621	CG	ARG	A	215	11.843	31.913	20.969	1.00	22.00
ATOM	1622	CD	ARG	A	215	11.231	33.041	21.727	1.00	23.87
ATOM	1623	NE	ARG	A	215	11.224	34.272	20.959	1.00	25.87
ATOM	1624	CZ	ARG	A	215	10.566	35.364	21.297	1.00	25.54
ATOM	1625	NH1	ARG	A	215	9.873	35.425	22.423	1.00	24.05
ATOM	1626	NH2	ARG	A	215	10.634	36.420	20.505	1.00	29.37
ATOM	1627	N	CYS	A	216	14.794	30.090	21.193	1.00	21.48
ATOM	1628	CA	CYS	A	216	16.123	30.355	21.668	1.00	21.40
ATOM	1629	C	CYS	A	216	16.533	31.723	21.146	1.00	21.86
ATOM	1630	O	CYS	A	216	16.532	31.952	19.948	1.00	22.82
ATOM	1631	CB	CYS	A	216	17.079	29.309	21.127	1.00	21.46
ATOM	1632	SG	CYS	A	216	16.689	27.562	21.469	1.00	22.63
ATOM	1633	N	ILE	A	217	16.878	32.630	22.045	1.00	21.29
ATOM	1634	CA	ILE	A	217	17.331	33.960	21.663	1.00	21.28
ATOM	1635	C	ILE	A	217	18.782	34.114	22.102	1.00	20.86
ATOM	1636	O	ILE	A	217	19.085	34.007	23.310	1.00	21.04
ATOM	1637	CB	ILE	A	217	16.456	35.025	22.327	1.00	21.80

O
C
C
C
C
N
C
C
O
C
C
C
N
N
C
C
O
C
C
C
C
C
O
N
C
C
C
O
C
C
C
N
N
N
C
C
C
O
C
S
N
C
C
O
C

ATOM	1638	CG1	ILE	A	217	14.978	34.807	21.955	1.00	22.90
ATOM	1639	CG2	ILE	A	217	16.874	36.414	21.861	1.00	22.72
ATOM	1640	CD1	ILE	A	217	14.031	35.695	22.693	1.00	24.99
ATOM	1641	N	LEU	A	218	19.666	34.346	21.128	1.00	20.25
ATOM	1642	CA	LEU	A	218	21.094	34.451	21.373	1.00	20.55
ATOM	1643	C	LEU	A	218	21.630	35.853	21.128	1.00	21.19
ATOM	1644	O	LEU	A	218	21.141	36.560	20.253	1.00	21.62
ATOM	1645	CB	LEU	A	218	21.874	33.493	20.467	1.00	20.31
ATOM	1646	CG	LEU	A	218	21.964	32.019	20.906	1.00	20.46
ATOM	1647	CD1	LEU	A	218	20.628	31.415	21.054	1.00	20.19
ATOM	1648	CD2	LEU	A	218	22.732	31.229	19.907	1.00	22.84
ATOM	1649	N	PHE	A	219	22.658	36.231	21.886	1.00	21.06
ATOM	1650	CA	PHE	A	219	23.331	37.498	21.682	1.00	21.62
ATOM	1651	C	PHE	A	219	24.842	37.244	21.611	1.00	21.87
ATOM	1652	O	PHE	A	219	25.390	36.490	22.416	1.00	21.88
ATOM	1653	CB	PHE	A	219	23.005	38.484	22.820	1.00	20.94
ATOM	1654	CG	PHE	A	219	21.541	38.617	23.106	1.00	21.13
ATOM	1655	CD1	PHE	A	219	20.907	37.779	23.995	1.00	21.92
ATOM	1656	CD2	PHE	A	219	20.785	39.589	22.476	1.00	24.21
ATOM	1657	CE1	PHE	A	219	19.563	37.915	24.250	1.00	21.95
ATOM	1658	CE2	PHE	A	219	19.442	39.724	22.735	1.00	22.06
ATOM	1659	CZ	PHE	A	219	18.840	38.904	23.621	1.00	23.11
ATOM	1660	N	PRO	A	220	25.512	37.877	20.662	1.00	22.38
ATOM	1661	CA	PRO	A	220	26.962	37.744	20.557	1.00	22.81
ATOM	1662	C	PRO	A	220	27.683	38.287	21.793	1.00	22.78
ATOM	1663	O	PRO	A	220	27.173	39.123	22.564	1.00	22.88
ATOM	1664	CB	PRO	A	220	27.335	38.561	19.317	1.00	22.81
ATOM	1665	CG	PRO	A	220	26.005	39.005	18.685	1.00	23.68
ATOM	1666	CD	PRO	A	220	24.940	38.792	19.663	1.00	22.86
ATOM	1667	N	PRO	A	221	28.885	37.774	21.999	1.00	22.84
ATOM	1668	CA	PRO	A	221	29.724	38.204	23.124	1.00	22.73
ATOM	1669	C	PRO	A	221	29.976	39.714	23.143	1.00	22.28
ATOM	1670	O	PRO	A	221	30.192	40.279	24.212	1.00	21.03
ATOM	1671	CB	PRO	A	221	31.030	37.455	22.892	1.00	22.78
ATOM	1672	CG	PRO	A	221	30.644	36.296	22.026	1.00	23.22
ATOM	1673	CD	PRO	A	221	29.521	36.733	21.174	1.00	22.58
ATOM	1674	N	ASP	A	222	29.897	40.378	21.993	1.00	22.60
ATOM	1675	CA	ASP	A	222	30.172	41.821	21.965	1.00	22.84
ATOM	1676	C	ASP	A	222	28.996	42.623	22.493	1.00	23.36
ATOM	1677	O	ASP	A	222	29.031	43.847	22.500	1.00	24.43
ATOM	1678	CB	ASP	A	222	30.638	42.308	20.579	1.00	22.47
ATOM	1679	CG	ASP	A	222	29.502	42.389	19.537	1.00	25.07
ATOM	1680	OD1	ASP	A	222	28.355	41.942	19.764	1.00	25.24
ATOM	1681	OD2	ASP	A	222	29.690	42.876	18.410	1.00	29.46
ATOM	1682	N	GLN	A	223	27.940	41.940	22.926	1.00	23.79
ATOM	1683	CA	GLN	A	223	26.826	42.624	23.565	1.00	23.85
ATOM	1684	C	GLN	A	223	26.942	42.570	25.114	1.00	23.37
ATOM	1685	O	GLN	A	223	25.986	42.875	25.850	1.00	23.78
ATOM	1686	CB	GLN	A	223	25.510	42.084	23.007	1.00	24.80
ATOM	1687	CG	GLN	A	223	25.217	42.672	21.550	1.00	28.46
ATOM	1688	CD	GLN	A	223	23.786	42.471	21.094	1.00	32.35
ATOM	1689	OE1	GLN	A	223	22.861	42.619	21.893	1.00	36.73
ATOM	1690	NE2	GLN	A	223	23.592	42.105	19.815	1.00	33.48
ATOM	1691	N	PHE	A	224	28.111	42.167	25.601	1.00	21.59
ATOM	1692	CA	PHE	A	224	28.394	42.195	27.035	1.00	21.84
ATOM	1693	C	PHE	A	224	27.888	43.509	27.706	1.00	21.56
ATOM	1694	O	PHE	A	224	27.233	43.455	28.738	1.00	20.27
ATOM	1695	CB	PHE	A	224	29.917	42.068	27.271	1.00	21.34
ATOM	1696	CG	PHE	A	224	30.309	41.943	28.732	1.00	22.22
ATOM	1697	CD1	PHE	A	224	30.449	43.071	29.540	1.00	21.80
ATOM	1698	CD2	PHE	A	224	30.583	40.723	29.279	1.00	20.92

ATOM	1699	CE1	PHE	A	224	30.842	42.945	30.868	1.00	22.58
ATOM	1700	CE2	PHE	A	224	30.979	40.593	30.587	1.00	22.14
ATOM	1701	CZ	PHE	A	224	31.087	41.725	31.394	1.00	22.16
ATOM	1702	N	GLU	A	225	28.190	44.665	27.106	1.00	22.03
ATOM	1703	CA	GLU	A	225	27.813	45.963	27.677	1.00	23.55
ATOM	1704	C	GLU	A	225	26.322	46.216	27.764	1.00	23.06
ATOM	1705	O	GLU	A	225	25.907	47.129	28.483	1.00	22.10
ATOM	1706	CB	GLU	A	225	28.396	47.111	26.856	1.00	24.92
ATOM	1707	CG	GLU	A	225	29.897	47.228	27.013	1.00	30.17
ATOM	1708	CD	GLU	A	225	30.386	48.632	27.280	1.00	36.52
ATOM	1709	OE1	GLU	A	225	29.901	49.319	28.242	1.00	40.52
ATOM	1710	OE2	GLU	A	225	31.310	49.028	26.532	1.00	41.63
ATOM	1711	N	CYS	A	226	25.532	45.456	27.003	1.00	22.07
ATOM	1712	CA	CYS	A	226	24.087	45.619	27.027	1.00	22.09
ATOM	1713	C	CYS	A	226	23.372	44.631	27.935	1.00	21.89
ATOM	1714	O	CYS	A	226	22.170	44.781	28.162	1.00	20.73
ATOM	1715	CB	CYS	A	226	23.523	45.416	25.630	1.00	22.22
ATOM	1716	SG	CYS	A	226	24.206	46.488	24.368	1.00	22.59
ATOM	1717	N	LEU	A	227	24.102	43.633	28.451	1.00	22.08
ATOM	1718	CA	LEU	A	227	23.467	42.529	29.169	1.00	22.08
ATOM	1719	C	LEU	A	227	23.771	42.393	30.657	1.00	21.99
ATOM	1720	O	LEU	A	227	23.118	41.611	31.373	1.00	22.28
ATOM	1721	CB	LEU	A	227	23.751	41.244	28.418	1.00	22.29
ATOM	1722	CG	LEU	A	227	22.874	41.185	27.158	1.00	23.54
ATOM	1723	CD1	LEU	A	227	23.428	40.186	26.144	1.00	23.51
ATOM	1724	CD2	LEU	A	227	21.423	40.819	27.544	1.00	24.49
ATOM	1725	N	TYR	A	228	24.763	43.140	31.109	1.00	20.87
ATOM	1726	CA	TYR	A	228	24.996	43.320	32.521	1.00	21.33
ATOM	1727	C	TYR	A	228	25.018	42.057	33.382	1.00	21.27
ATOM	1728	O	TYR	A	228	24.205	41.920	34.301	1.00	21.72
ATOM	1729	CB	TYR	A	228	23.957	44.311	33.077	1.00	20.92
ATOM	1730	CG	TYR	A	228	23.949	45.649	32.339	1.00	20.91
ATOM	1731	CD1	TYR	A	228	24.768	46.677	32.739	1.00	18.87
ATOM	1732	CD2	TYR	A	228	23.122	45.862	31.232	1.00	20.51
ATOM	1733	CE1	TYR	A	228	24.780	47.912	32.076	1.00	20.36
ATOM	1734	CE2	TYR	A	228	23.117	47.074	30.561	1.00	21.07
ATOM	1735	CZ	TYR	A	228	23.931	48.113	31.000	1.00	20.84
ATOM	1736	OH	TYR	A	228	23.947	49.325	30.336	1.00	19.16
ATOM	1737	N	PRO	A	229	25.985	41.175	33.141	1.00	20.77
ATOM	1738	CA	PRO	A	229	26.128	39.971	33.966	1.00	21.05
ATOM	1739	C	PRO	A	229	26.428	40.332	35.408	1.00	20.08
ATOM	1740	O	PRO	A	229	27.021	41.379	35.675	1.00	20.20
ATOM	1741	CB	PRO	A	229	27.360	39.271	33.363	1.00	20.51
ATOM	1742	CG	PRO	A	229	28.084	40.340	32.651	1.00	21.64
ATOM	1743	CD	PRO	A	229	27.025	41.241	32.105	1.00	20.97
ATOM	1744	N	TYR	A	230	25.988	39.490	36.326	1.00	19.73
ATOM	1745	CA	TYR	A	230	26.282	39.656	37.724	1.00	18.80
ATOM	1746	C	TYR	A	230	27.809	39.657	37.947	1.00	18.61
ATOM	1747	O	TYR	A	230	28.575	39.250	37.094	1.00	18.26
ATOM	1748	CB	TYR	A	230	25.669	38.507	38.520	1.00	18.86
ATOM	1749	CG	TYR	A	230	24.172	38.597	38.725	1.00	19.28
ATOM	1750	CD1	TYR	A	230	23.297	38.110	37.760	1.00	18.53
ATOM	1751	CD2	TYR	A	230	23.629	39.172	39.885	1.00	19.38
ATOM	1752	CE1	TYR	A	230	21.932	38.161	37.935	1.00	18.86
ATOM	1753	CE2	TYR	A	230	22.234	39.222	40.076	1.00	19.53
ATOM	1754	CZ	TYR	A	230	21.404	38.719	39.073	1.00	20.93
ATOM	1755	OH	TYR	A	230	20.034	38.771	39.176	1.00	23.69
ATOM	1756	N	PRO	A	231	28.251	40.137	39.098	1.00	18.40
ATOM	1757	CA	PRO	A	231	29.651	39.983	39.476	1.00	18.58
ATOM	1758	C	PRO	A	231	30.070	38.507	39.399	1.00	19.41
ATOM	1759	O	PRO	A	231	29.240	37.624	39.654	1.00	18.89

C
C
C
N
C
C
O
C
C
O
O
N
C
C
O
C
S
N
C
C
O
C
C
C
N
C
C
O
C
C
O
N
C
C
O
C
C
C
N
C
C
O
C
C
C
O
N
C
C
O
N
C
C
O

ATOM	1760	CB	PRO A 231	29.675	40.477	40.931	1.00	18.95
ATOM	1761	CG	PRO A 231	28.542	41.507	41.001	1.00	19.02
ATOM	1762	CD	PRO A 231	27.460	40.868	40.105	1.00	18.02
ATOM	1763	N	VAL A 232	31.335	38.248	39.071	1.00	19.55
ATOM	1764	CA	VAL A 232	31.826	36.884	38.969	1.00	20.65
ATOM	1765	C	VAL A 232	31.640	36.034	40.236	1.00	20.79
ATOM	1766	O	VAL A 232	31.379	34.838	40.145	1.00	21.82
ATOM	1767	CB	VAL A 232	33.318	36.869	38.545	1.00	20.71
ATOM	1768	CG1	VAL A 232	33.945	35.524	38.807	1.00	21.96
ATOM	1769	CG2	VAL A 232	33.452	37.225	37.075	1.00	20.83
ATOM	1770	N	HIS A 233	31.770	36.639	41.409	1.00	20.87
ATOM	1771	CA	HIS A 233	31.643	35.907	42.661	1.00	20.70
ATOM	1772	C	HIS A 233	30.203	35.739	43.168	1.00	20.59
ATOM	1773	O	HIS A 233	29.940	35.010	44.119	1.00	20.49
ATOM	1774	CB	HIS A 233	32.482	36.593	43.706	1.00	20.63
ATOM	1775	CG	HIS A 233	33.948	36.529	43.426	1.00	21.92
ATOM	1776	ND1	HIS A 233	34.659	37.595	42.913	1.00	22.17
ATOM	1777	CD2	HIS A 233	34.843	35.527	43.610	1.00	22.03
ATOM	1778	CE1	HIS A 233	35.929	37.249	42.793	1.00	23.69
ATOM	1779	NE2	HIS A 233	36.066	35.998	43.203	1.00	23.04
ATOM	1780	N	HIS A 234	29.273	36.433	42.543	1.00	20.53
ATOM	1781	CA	HIS A 234	27.869	36.251	42.857	1.00	20.60
ATOM	1782	C	HIS A 234	27.388	34.895	42.306	1.00	20.45
ATOM	1783	O	HIS A 234	27.873	34.428	41.291	1.00	19.86
ATOM	1784	CB	HIS A 234	27.088	37.359	42.206	1.00	20.65
ATOM	1785	CG	HIS A 234	25.661	37.441	42.625	1.00	20.68
ATOM	1786	ND1	HIS A 234	24.667	36.700	42.021	1.00	21.46
ATOM	1787	CD2	HIS A 234	25.044	38.249	43.518	1.00	20.20
ATOM	1788	CE1	HIS A 234	23.504	37.013	42.563	1.00	21.99
ATOM	1789	NE2	HIS A 234	23.707	37.951	43.473	1.00	21.29
ATOM	1790	N	PRO A 235	26.463	34.261	43.007	1.00	20.40
ATOM	1791	CA	PRO A 235	25.888	32.993	42.572	1.00	20.85
ATOM	1792	C	PRO A 235	25.360	33.021	41.152	1.00	21.20
ATOM	1793	O	PRO A 235	25.437	31.984	40.502	1.00	20.43
ATOM	1794	CB	PRO A 235	24.756	32.771	43.575	1.00	21.58
ATOM	1795	CG	PRO A 235	25.273	33.452	44.826	1.00	20.78
ATOM	1796	CD	PRO A 235	25.941	34.685	44.317	1.00	20.56
ATOM	1797	N	CYS A 236	24.905	34.173	40.663	1.00	20.46
ATOM	1798	CA	CYS A 236	24.379	34.235	39.308	1.00	20.63
ATOM	1799	C	CYS A 236	25.430	34.707	38.301	1.00	20.58
ATOM	1800	O	CYS A 236	25.110	35.283	37.258	1.00	21.60
ATOM	1801	CB	CYS A 236	23.114	35.084	39.268	1.00	20.06
ATOM	1802	SG	CYS A 236	21.824	34.377	40.326	1.00	21.70
ATOM	1803	N	ASP A 237	26.693	34.474	38.629	1.00	20.17
ATOM	1804	CA	ASP A 237	27.785	34.632	37.679	1.00	19.98
ATOM	1805	C	ASP A 237	27.381	34.050	36.313	1.00	19.79

C
C
O
N
C
C
O
C
C
N
C
C
O
C
C
N
C
C
N
N
C
C
O
C
C
N
C
C
N
N
C
C
C
C
C
C
N
C
C
C
S
N
C
C
C
O
C
C
O
N
C
C
C
C
C
N
C
N

ATOM	1821	NH2	ARG	A	238	27.399	28.949	35.036	1.00	27.47
ATOM	1822	N	GLN	A	239	25.021	35.028	34.375	1.00	20.56
ATOM	1823	CA	GLN	A	239	23.641	35.334	34.014	1.00	21.11
ATOM	1824	C	GLN	A	239	23.459	36.865	33.920	1.00	20.84
ATOM	1825	O	GLN	A	239	24.145	37.624	34.610	1.00	21.25
ATOM	1826	CB	GLN	A	239	22.646	34.776	35.027	1.00	20.14
ATOM	1827	CG	GLN	A	239	23.052	33.505	35.726	1.00	22.87
ATOM	1828	CD	GLN	A	239	23.281	32.320	34.807	1.00	23.97
ATOM	1829	OE1	GLN	A	239	22.367	31.842	34.132	1.00	23.63
ATOM	1830	NE2	GLN	A	239	24.502	31.834	34.796	1.00	24.34
ATOM	1831	N	SER	A	240	22.541	37.307	33.071	1.00	20.31
ATOM	1832	CA	SER	A	240	22.246	38.725	32.926	1.00	19.92
ATOM	1833	C	SER	A	240	21.448	39.218	34.119	1.00	20.07
ATOM	1834	O	SER	A	240	20.538	38.533	34.564	1.00	19.79
ATOM	1835	CB	SER	A	240	21.384	38.953	31.700	1.00	19.45
ATOM	1836	OG	SER	A	240	21.018	40.312	31.564	1.00	20.53
ATOM	1837	N	GLN	A	241	21.763	40.408	34.623	1.00	19.46
ATOM	1838	CA	GLN	A	241	20.962	40.988	35.701	1.00	20.32
ATOM	1839	C	GLN	A	241	19.657	41.569	35.175	1.00	20.51
ATOM	1840	O	GLN	A	241	18.808	41.956	35.959	1.00	21.06
ATOM	1841	CB	GLN	A	241	21.697	42.141	36.417	1.00	20.15
ATOM	1842	CG	GLN	A	241	22.910	41.749	37.230	1.00	21.41
ATOM	1843	CD	GLN	A	241	23.690	42.980	37.695	1.00	22.76
ATOM	1844	OE1	GLN	A	241	23.455	43.481	38.775	1.00	24.42
ATOM	1845	NE2	GLN	A	241	24.585	43.465	36.871	1.00	21.93
ATOM	1846	N	VAL	A	242	19.489	41.678	33.862	1.00	20.66
ATOM	1847	CA	VAL	A	242	18.309	42.355	33.372	1.00	20.76
ATOM	1848	C	VAL	A	242	17.089	41.465	33.391	1.00	21.26
ATOM	1849	O	VAL	A	242	17.119	40.352	32.871	1.00	21.01
ATOM	1850	CB	VAL	A	242	18.476	42.813	31.909	1.00	20.97
ATOM	1851	CG1	VAL	A	242	17.194	43.502	31.431	1.00	21.21
ATOM	1852	CG2	VAL	A	242	19.687	43.704	31.716	1.00	18.95
ATOM	1853	N	ASP	A	243	16.001	41.960	33.958	1.00	21.16
ATOM	1854	CA	ASP	A	243	14.730	41.223	33.907	1.00	21.36
ATOM	1855	C	ASP	A	243	14.091	41.523	32.552	1.00	21.20
ATOM	1856	O	ASP	A	243	13.566	42.631	32.310	1.00	19.97
ATOM	1857	CB	ASP	A	243	13.840	41.682	35.054	1.00	21.65
ATOM	1858	CG	ASP	A	243	12.474	41.051	35.037	1.00	22.45
ATOM	1859	OD1	ASP	A	243	12.122	40.338	34.070	1.00	25.16
ATOM	1860	OD2	ASP	A	243	11.671	41.232	35.978	1.00	25.29
ATOM	1861	N	PHE	A	244	14.182	40.554	31.650	1.00	21.28
ATOM	1862	CA	PHE	A	244	13.687	40.733	30.291	1.00	21.65
ATOM	1863	C	PHE	A	244	12.202	41.074	30.274	1.00	22.36
ATOM	1864	O	PHE	A	244	11.733	41.653	29.304	1.00	22.09
ATOM	1865	CB	PHE	A	244	13.915	39.478	29.442	1.00	22.22
ATOM	1866	CG	PHE	A	244	15.284	39.370	28.815	1.00	21.43
ATOM	1867	CD1	PHE	A	244	16.414	39.800	29.465	1.00	20.12
ATOM	1868	CD2	PHE	A	244	15.422	38.778	27.575	1.00	21.05
ATOM	1869	CE1	PHE	A	244	17.655	39.671	28.873	1.00	20.83
ATOM	1870	CE2	PHE	A	244	16.645	38.643	26.993	1.00	21.47
ATOM	1871	CZ	PHE	A	244	17.767	39.104	27.633	1.00	20.74
ATOM	1872	N	ASP	A	245	11.464	40.696	31.322	1.00	23.26
ATOM	1873	CA	ASP	A	245	10.035	40.973	31.393	1.00	24.31
ATOM	1874	C	ASP	A	245	9.756	42.401	31.877	1.00	25.05
ATOM	1875	O	ASP	A	245	8.679	42.916	31.646	1.00	24.27
ATOM	1876	CB	ASP	A	245	9.333	40.004	32.348	1.00	24.79
ATOM	1877	CG	ASP	A	245	9.397	38.571	31.887	1.00	25.46
ATOM	1878	OD1	ASP	A	245	9.409	38.342	30.663	1.00	27.47
ATOM	1879	OD2	ASP	A	245	9.445	37.610	32.678	1.00	25.22
ATOM	1880	N	ASN	A	246	10.702	43.006	32.589	1.00	25.30
ATOM	1881	CA	ASN	A	246	10.531	44.365	33.099	1.00	26.37

ATOM	1882	C	ASN	A	246	11.876	45.054	33.204	1.00	25.61
ATOM	1883	O	ASN	A	246	12.436	45.192	34.282	1.00	26.05
ATOM	1884	CB	ASN	A	246	9.871	44.360	34.472	1.00	26.89
ATOM	1885	CG	ASN	A	246	9.545	45.770	34.958	1.00	30.93
ATOM	1886	OD1	ASN	A	246	9.353	46.701	34.150	1.00	35.10
ATOM	1887	ND2	ASN	A	246	9.493	45.943	36.281	1.00	35.04
ATOM	1888	N	PRO	A	247	12.415	45.450	32.068	1.00	24.71
ATOM	1889	CA	PRO	A	247	13.775	45.979	32.026	1.00	24.24
ATOM	1890	C	PRO	A	247	13.909	47.352	32.684	1.00	24.48
ATOM	1891	O	PRO	A	247	13.189	48.296	32.374	1.00	23.54
ATOM	1892	CB	PRO	A	247	14.109	46.036	30.538	1.00	24.23
ATOM	1893	CG	PRO	A	247	12.900	45.524	29.803	1.00	25.13
ATOM	1894	CD	PRO	A	247	11.769	45.385	30.752	1.00	24.61
ATOM	1895	N	ASP	A	248	14.867	47.437	33.593	1.00	24.04
ATOM	1896	CA	ASP	A	248	15.140	48.656	34.285	1.00	24.53
ATOM	1897	C	ASP	A	248	16.206	49.440	33.496	1.00	24.15
ATOM	1898	O	ASP	A	248	17.410	49.221	33.629	1.00	23.11
ATOM	1899	CB	ASP	A	248	15.609	48.312	35.689	1.00	24.75
ATOM	1900	CG	ASP	A	248	15.731	49.512	36.554	1.00	26.08
ATOM	1901	OD1	ASP	A	248	15.956	50.616	36.009	1.00	27.08
ATOM	1902	OD2	ASP	A	248	15.623	49.439	37.797	1.00	30.33
ATOM	1903	N	TYR	A	249	15.735	50.329	32.642	1.00	24.42
ATOM	1904	CA	TYR	A	249	16.612	51.129	31.807	1.00	25.30
ATOM	1905	C	TYR	A	249	17.462	52.136	32.589	1.00	26.13
ATOM	1906	O	TYR	A	249	18.495	52.584	32.083	1.00	26.46
ATOM	1907	CB	TYR	A	249	15.796	51.826	30.713	1.00	25.41
ATOM	1908	CG	TYR	A	249	15.119	50.853	29.768	1.00	22.90
ATOM	1909	CD1	TYR	A	249	15.844	49.873	29.130	1.00	22.91
ATOM	1910	CD2	TYR	A	249	13.760	50.923	29.519	1.00	22.21
ATOM	1911	CE1	TYR	A	249	15.238	48.964	28.271	1.00	22.19
ATOM	1912	CE2	TYR	A	249	13.144	50.023	28.660	1.00	21.56
ATOM	1913	CZ	TYR	A	249	13.895	49.044	28.046	1.00	21.23
ATOM	1914	OH	TYR	A	249	13.304	48.163	27.173	1.00	23.38
ATOM	1915	N	GLU	A	250	17.071	52.465	33.822	1.00	26.65
ATOM	1916	CA	GLU	A	250	17.894	53.365	34.640	1.00	27.47
ATOM	1917	C	GLU	A	250	19.161	52.651	35.086	1.00	26.52
ATOM	1918	O	GLU	A	250	20.238	53.224	35.086	1.00	27.17
ATOM	1919	CB	GLU	A	250	17.133	53.908	35.866	1.00	27.89
ATOM	1920	CG	GLU	A	250	15.880	54.684	35.485	1.00	32.57
ATOM	1921	CD	GLU	A	250	15.258	55.459	36.631	1.00	37.09
ATOM	1922	OE1	GLU	A	250	15.809	55.491	37.753	1.00	42.28
ATOM	1923	OE2	GLU	A	250	14.198	56.055	36.399	1.00	42.29
ATOM	1924	N	ARG	A	251	19.042	51.399	35.490	1.00	25.57
ATOM	1925	CA	ARG	A	251	20.224	50.657	35.900	1.00	24.72
ATOM	1926	C	ARG	A	251	20.955	50.069	34.718	1.00	23.56
ATOM	1927	O	ARG	A	251	22.150	49.919	34.769	1.00	22.92
ATOM	1928	CB	ARG	A	251	19.845	49.520	36.845	1.00	25.64
ATOM	1929	CG	ARG	A	251	19.435	49.957	38.255	1.00	26.79
ATOM	1930	CD	ARG	A	251	18.858	48.815	39.123	1.00	30.01
ATOM	1931	NE	ARG	A	251	19.900	47.875	39.540	1.00	31.06
ATOM	1932	CZ	ARG	A	251	19.709	46.579	39.766	1.00	32.72
ATOM	1933	NH1	ARG	A	251	18.508	46.029	39.628	1.00	33.10
ATOM	1934	NH2	ARG	A	251	20.734	45.828	40.135	1.00	33.44
ATOM	1935	N	PHE	A	252	20.234	49.725	33.652	1.00	22.66
ATOM	1936	CA	PHE	A	252	20.826	49.019	32.532	1.00	21.99
ATOM	1937	C	PHE	A	252	20.521	49.719	31.198	1.00	21.71
ATOM	1938	O	PHE	A	252	19.900	49.160	30.289	1.00	21.32
ATOM	1939	CB	PHE	A	252	20.261	47.610	32.519	1.00	22.03
ATOM	1940	CG	PHE	A	252	20.292	46.915	33.868	1.00	21.75
ATOM	1941	CD1	PHE	A	252	21.484	46.729	34.556	1.00	20.90
ATOM	1942	CD2	PHE	A	252	19.124	46.402	34.423	1.00	21.94

C
O
C
C
O
N
N
C
C
O
C
C
C
N
C
C
O
C
C
O
C
C
C
O
N
C
C
O
C
C
C
O
N
C
C
O
C
C
C
N
C
N
N
N
C
C
C
C
C
C

ATOM	1943	CE1	PHE	A	252	21.506	46.040	35.785	1.00	20.51
ATOM	1944	CE2	PHE	A	252	19.136	45.730	35.662	1.00	21.77
ATOM	1945	CZ	PHE	A	252	20.320	45.553	36.333	1.00	21.41
ATOM	1946	N	PRO	A	253	20.989	50.944	31.071	1.00	20.88
ATOM	1947	CA	PRO	A	253	20.615	51.754	29.909	1.00	20.51
ATOM	1948	C	PRO	A	253	20.956	51.077	28.578	1.00	20.47
ATOM	1949	O	PRO	A	253	20.153	51.159	27.656	1.00	20.01
ATOM	1950	CB	PRO	A	253	21.371	53.077	30.130	1.00	20.00
ATOM	1951	CG	PRO	A	253	22.538	52.697	31.136	1.00	20.46
ATOM	1952	CD	PRO	A	253	21.921	51.641	31.994	1.00	20.89
ATOM	1953	N	ASN	A	254	22.088	50.393	28.455	1.00	20.68
ATOM	1954	CA	ASN	A	254	22.401	49.805	27.160	1.00	20.72
ATOM	1955	C	ASN	A	254	21.537	48.601	26.790	1.00	20.29
ATOM	1956	O	ASN	A	254	21.644	48.068	25.688	1.00	20.03
ATOM	1957	CB	ASN	A	254	23.882	49.463	27.033	1.00	20.89
ATOM	1958	CG	ASN	A	254	24.736	50.698	26.899	1.00	22.24
ATOM	1959	OD1	ASN	A	254	25.532	50.999	27.785	1.00	24.48
ATOM	1960	ND2	ASN	A	254	24.557	51.446	25.791	1.00	22.23
ATOM	1961	N	PHE	A	255	20.678	48.160	27.690	1.00	20.01
ATOM	1962	CA	PHE	A	255	19.781	47.086	27.305	1.00	20.02
ATOM	1963	C	PHE	A	255	18.855	47.599	26.185	1.00	20.33
ATOM	1964	O	PHE	A	255	18.224	46.818	25.472	1.00	19.82
ATOM	1965	CB	PHE	A	255	18.970	46.552	28.484	1.00	19.63
ATOM	1966	CG	PHE	A	255	18.239	45.318	28.145	1.00	18.96
ATOM	1967	CD1	PHE	A	255	18.928	44.149	27.930	1.00	19.50
ATOM	1968	CD2	PHE	A	255	16.889	45.335	27.942	1.00	17.83
ATOM	1969	CE1	PHE	A	255	18.267	42.997	27.575	1.00	20.08
ATOM	1970	CE2	PHE	A	255	16.230	44.206	27.561	1.00	17.77
ATOM	1971	CZ	PHE	A	255	16.912	43.036	27.381	1.00	21.57
ATOM	1972	N	GLN	A	256	18.788	48.919	26.035	1.00	20.15
ATOM	1973	CA	GLN	A	256	17.973	49.533	24.986	1.00	20.63
ATOM	1974	C	GLN	A	256	18.594	49.343	23.591	1.00	20.12
ATOM	1975	O	GLN	A	256	17.955	49.612	22.592	1.00	19.87
ATOM	1976	CB	GLN	A	256	17.787	51.036	25.272	1.00	20.67
ATOM	1977	CG	GLN	A	256	16.744	51.306	26.362	1.00	22.56
ATOM	1978	CD	GLN	A	256	16.747	52.732	26.888	1.00	22.75
ATOM	1979	OE1	GLN	A	256	15.727	53.420	26.816	1.00	24.69
ATOM	1980	NE2	GLN	A	256	17.873	53.172	27.419	1.00	23.90
ATOM	1981	N	ASN	A	257	19.850	48.915	23.553	1.00	20.07
ATOM	1982	CA	ASN	A	257	20.583	48.700	22.310	1.00	20.66
ATOM	1983	C	ASN	A	257	20.778	47.233	21.963	1.00	21.21
ATOM	1984	O	ASN	A	257	21.446	46.921	20.979	1.00	21.28
ATOM	1985	CB	ASN	A	257	21.975	49.300	22.420	1.00	20.31
ATOM	1986	CG	ASN	A	257	21.937	50.757	22.776	1.00	20.83
ATOM	1987	OD1	ASN	A	257	22.395	51.141	23.850	1.00	20.15
ATOM	1988	ND2	ASN	A	257	21.366	51.577	21.889	1.00	16.99
ATOM	1989	N	VAL	A	258	20.218	46.333	22.758	1.00	21.91
ATOM	1990	CA	VAL	A	258	20.440	44.908	22.532	1.00	23.55
ATOM	1991	C	VAL	A	258	19.720	44.400	21.284	1.00	24.29
ATOM	1992	O	VAL	A	258	18.600	44.845	20.947	1.00	24.28
ATOM	1993	CB	VAL	A	258	20.026	44.110	23.772	1.00	24.18
ATOM	1994	CG1	VAL	A	258	18.529	43.868	23.785	1.00	23.69
ATOM	1995	CG2	VAL	A	258	20.761	42.851	23.826	1.00	28.07
ATOM	1996	N	VAL	A	259	20.381	43.506	20.562	1.00	24.48
ATOM	1997	CA	VAL	A	259	19.787	42.922	19.364	1.00	25.21
ATOM	1998	C	VAL	A	259	20.104	41.427	19.331	1.00	25.19
ATOM	1999	O	VAL	A	259	21.260	41.046	19.333	1.00	25.58
ATOM	2000	CB	VAL	A	259	20.345	43.551	18.094	1.00	25.37
ATOM	2001	CG1	VAL	A	259	19.726	42.893	16.882	1.00	27.15
ATOM	2002	CG2	VAL	A	259	20.074	45.059	18.066	1.00	25.97
ATOM	2003	N	GLY	A	260	19.068	40.603	19.308	1.00	25.20

[illegible]

ATOM	2004	CA	GLY	A	260	19.214	39.161	19.312	1.00	25.52
ATOM	2005	C	GLY	A	260	19.108	38.433	17.961	1.00	25.26
ATOM	2006	O	GLY	A	260	18.762	39.016	16.914	1.00	24.66
ATOM	2007	N	TYR	A	261	19.484	37.160	18.015	1.00	24.49
ATOM	2008	CA	TYR	A	261	19.379	36.217	16.925	1.00	25.05
ATOM	2009	C	TYR	A	261	18.435	35.145	17.475	1.00	24.39
ATOM	2010	O	TYR	A	261	18.671	34.620	18.552	1.00	23.82
ATOM	2011	CB	TYR	A	261	20.730	35.574	16.627	1.00	25.82
ATOM	2012	CG	TYR	A	261	21.748	36.476	15.965	1.00	28.98
ATOM	2013	CD1	TYR	A	261	21.804	36.598	14.585	1.00	35.41
ATOM	2014	CD2	TYR	A	261	22.649	37.194	16.712	1.00	29.78
ATOM	2015	CE1	TYR	A	261	22.737	37.429	13.976	1.00	35.73
ATOM	2016	CE2	TYR	A	261	23.592	37.994	16.126	1.00	32.54
ATOM	2017	CZ	TYR	A	261	23.643	38.116	14.768	1.00	35.98
ATOM	2018	OH	TYR	A	261	24.580	38.959	14.202	1.00	38.02
ATOM	2019	N	GLU	A	262	17.353	34.823	16.790	1.00	24.04
ATOM	2020	CA	GLU	A	262	16.432	33.860	17.387	1.00	23.96
ATOM	2021	C	GLU	A	262	15.992	32.773	16.435	1.00	23.90
ATOM	2022	O	GLU	A	262	16.116	32.890	15.213	1.00	23.56
ATOM	2023	CB	GLU	A	262	15.224	34.553	18.020	1.00	23.70
ATOM	2024	CG	GLU	A	262	14.029	34.782	17.127	1.00	24.33
ATOM	2025	CD	GLU	A	262	12.829	35.382	17.870	1.00	26.49
ATOM	2026	OE1	GLU	A	262	12.753	36.611	18.029	1.00	27.38
ATOM	2027	OE2	GLU	A	262	11.946	34.631	18.306	1.00	28.26
ATOM	2028	N	THR	A	263	15.485	31.713	17.034	1.00	23.68
ATOM	2029	CA	THR	A	263	14.974	30.605	16.279	1.00	24.40
ATOM	2030	C	THR	A	263	14.091	29.786	17.193	1.00	24.29
ATOM	2031	O	THR	A	263	14.131	29.936	18.423	1.00	24.53
ATOM	2032	CB	THR	A	263	16.148	29.762	15.751	1.00	24.52
ATOM	2033	OG1	THR	A	263	15.684	28.839	14.762	1.00	24.43
ATOM	2034	CG2	THR	A	263	16.737	28.876	16.854	1.00	24.33
ATOM	2035	N	VAL	A	264	13.263	28.954	16.593	1.00	23.62
ATOM	2036	CA	VAL	A	264	12.500	28.019	17.370	1.00	24.01
ATOM	2037	C	VAL	A	264	12.936	26.627	16.935	1.00	24.53
ATOM	2038	O	VAL	A	264	12.879	26.291	15.768	1.00	24.27
ATOM	2039	CB	VAL	A	264	11.008	28.194	17.177	1.00	24.69
ATOM	2040	CG1	VAL	A	264	10.256	26.958	17.712	1.00	23.69
ATOM	2041	CG2	VAL	A	264	10.520	29.526	17.857	1.00	23.81
ATOM	2042	N	VAL	A	265	13.450	25.844	17.872	1.00	24.91
ATOM	2043	CA	VAL	A	265	13.833	24.484	17.540	1.00	24.93
ATOM	2044	C	VAL	A	265	12.792	23.470	17.967	1.00	24.20
ATOM	2045	O	VAL	A	265	12.167	23.617	19.018	1.00	24.01

ATOM	2065	CA	ASP	A	269	16.849	20.141	15.364	1.00	19.78
ATOM	2066	C	ASP	A	269	17.991	20.162	16.395	1.00	19.63
ATOM	2067	O	ASP	A	269	17.829	19.729	17.551	1.00	18.28
ATOM	2068	CB	ASP	A	269	16.069	21.436	15.532	1.00	20.34
ATOM	2069	CG	ASP	A	269	14.976	21.636	14.500	1.00	20.66
ATOM	2070	OD1	ASP	A	269	14.977	21.015	13.419	1.00	22.15
ATOM	2071	OD2	ASP	A	269	14.072	22.454	14.703	1.00	23.30
ATOM	2072	N	VAL	A	270	19.129	20.681	15.953	1.00	19.76
ATOM	2073	CA	VAL	A	270	20.304	20.879	16.782	1.00	20.24
ATOM	2074	C	VAL	A	270	20.793	22.318	16.601	1.00	19.84
ATOM	2075	O	VAL	A	270	21.052	22.754	15.487	1.00	19.90
ATOM	2076	CB	VAL	A	270	21.421	19.942	16.377	1.00	20.08
ATOM	2077	CG1	VAL	A	270	22.676	20.288	17.121	1.00	21.38
ATOM	2078	CG2	VAL	A	270	21.017	18.521	16.671	1.00	21.15
ATOM	2079	N	LEU	A	271	20.885	23.056	17.699	1.00	19.86
ATOM	2080	CA	LEU	A	271	21.341	24.427	17.656	1.00	20.03
ATOM	2081	C	LEU	A	271	22.757	24.465	18.202	1.00	20.23
ATOM	2082	O	LEU	A	271	23.032	23.946	19.276	1.00	19.73
ATOM	2083	CB	LEU	A	271	20.429	25.328	18.465	1.00	20.59
ATOM	2084	CG	LEU	A	271	20.934	26.742	18.724	1.00	20.93
ATOM	2085	CD1	LEU	A	271	21.011	27.562	17.431	1.00	20.35
ATOM	2086	CD2	LEU	A	271	20.039	27.428	19.728	1.00	22.31
ATOM	2087	N	TYR	A	272	23.669	25.024	17.416	1.00	20.58
ATOM	2088	CA	TYR	A	272	25.010	25.229	17.883	1.00	20.58
ATOM	2089	C	TYR	A	272	24.985	26.535	18.653	1.00	20.69
ATOM	2090	O	TYR	A	272	24.692	27.576	18.089	1.00	21.54
ATOM	2091	CB	TYR	A	272	26.004	25.312	16.728	1.00	20.30
ATOM	2092	CG	TYR	A	272	27.381	25.776	17.158	1.00	21.24
ATOM	2093	CD1	TYR	A	272	27.997	25.250	18.275	1.00	21.70
ATOM	2094	CD2	TYR	A	272	28.039	26.788	16.472	1.00	22.21
ATOM	2095	CE1	TYR	A	272	29.232	25.692	18.674	1.00	21.69
ATOM	2096	CE2	TYR	A	272	29.275	27.215	16.857	1.00	22.99
ATOM	2097	CZ	TYR	A	272	29.868	26.672	17.961	1.00	22.01
ATOM	2098	OH	TYR	A	272	31.109	27.111	18.352	1.00	22.77
ATOM	2099	N	ILE	A	273	25.216	26.454	19.951	1.00	21.22
ATOM	2100	CA	ILE	A	273	25.362	27.629	20.805	1.00	21.49
ATOM	2101	C	ILE	A	273	26.852	27.809	21.117	1.00	22.00
ATOM	2102	O	ILE	A	273	27.413	27.112	21.962	1.00	22.55
ATOM	2103	CB	ILE	A	273	24.607	27.471	22.073	1.00	21.14
ATOM	2104	CG1	ILE	A	273	23.148	27.132	21.776	1.00	21.68
ATOM	2105	CG2	ILE	A	273	24.661	28.791	22.854	1.00	21.75
ATOM	2106	CD1	ILE	A	273	22.318	26.972	23.026	1.00	21.44
ATOM	2107	N	PRO	A	274					

ATOM	2126	CB	TYR	A	276	31.673	33.594	24.934	1.00	23.42
ATOM	2127	CG	TYR	A	276	32.857	33.283	25.796	1.00	28.51
ATOM	2128	CD1	TYR	A	276	32.969	33.799	27.072	1.00	33.00
ATOM	2129	CD2	TYR	A	276	33.876	32.460	25.321	1.00	34.41
ATOM	2130	CE1	TYR	A	276	34.066	33.490	27.877	1.00	36.36
ATOM	2131	CE2	TYR	A	276	34.968	32.151	26.102	1.00	36.45
ATOM	2132	CZ	TYR	A	276	35.059	32.651	27.371	1.00	37.63
ATOM	2133	OH	TYR	A	276	36.192	32.314	28.077	1.00	40.88
ATOM	2134	N	TRP	A	277	28.455	32.878	23.906	1.00	20.75
ATOM	2135	CA	TRP	A	277	27.272	33.528	23.360	1.00	20.85
ATOM	2136	C	TRP	A	277	26.173	33.482	24.372	1.00	20.33
ATOM	2137	O	TRP	A	277	25.907	32.431	24.949	1.00	20.72
ATOM	2138	CB	TRP	A	277	26.791	32.874	22.068	1.00	20.13
ATOM	2139	CG	TRP	A	277	27.552	33.323	20.879	1.00	19.28
ATOM	2140	CD1	TRP	A	277	28.887	33.175	20.673	1.00	21.27
ATOM	2141	CD2	TRP	A	277	27.042	33.995	19.730	1.00	20.60
ATOM	2142	NE1	TRP	A	277	29.246	33.721	19.467	1.00	21.05
ATOM	2143	CE2	TRP	A	277	28.128	34.232	18.864	1.00	21.04
ATOM	2144	CE3	TRP	A	277	25.780	34.456	19.352	1.00	21.08
ATOM	2145	CZ2	TRP	A	277	27.989	34.893	17.643	1.00	20.38
ATOM	2146	CZ3	TRP	A	277	25.641	35.110	18.122	1.00	20.48
ATOM	2147	CH2	TRP	A	277	26.736	35.306	17.288	1.00	21.03
ATOM	2148	N	TRP	A	278	25.572	34.632	24.629	1.00	19.39
ATOM	2149	CA	TRP	A	278	24.430	34.711	25.521	1.00	19.80
ATOM	2150	C	TRP	A	278	23.243	33.927	24.927	1.00	20.18
ATOM	2151	O	TRP	A	278	23.019	33.939	23.732	1.00	21.04
ATOM	2152	CB	TRP	A	278	23.989	36.168	25.646	1.00	20.39
ATOM	2153	CG	TRP	A	278	24.979	37.031	26.300	1.00	21.20
ATOM	2154	CD1	TRP	A	278	26.022	37.672	25.720	1.00	21.06
ATOM	2155	CD2	TRP	A	278	25.056	37.301	27.691	1.00	20.38
ATOM	2156	NE1	TRP	A	278	26.725	38.369	26.668	1.00	22.44
ATOM	2157	CE2	TRP	A	278	26.139	38.155	27.891	1.00	21.86
ATOM	2158	CE3	TRP	A	278	24.282	36.937	28.789	1.00	21.09
ATOM	2159	CZ2	TRP	A	278	26.480	38.640	29.147	1.00	21.13
ATOM	2160	CZ3	TRP	A	278	24.623	37.412	30.019	1.00	23.38
ATOM	2161	CH2	TRP	A	278	25.704	38.258	30.194	1.00	21.06
ATOM	2162	N	HIS	A	279	22.465	33.278	25.761	1.00	20.76
ATOM	2163	CA	HIS	A	279	21.262	32.646	25.294	1.00	21.56
ATOM	2164	C	HIS	A	279	20.198	32.659	26.361	1.00	21.24
ATOM	2165	O	HIS	A	279	20.475	32.456	27.540	1.00	21.63
ATOM	2166	CB	HIS	A	279	21.501	31.203	24.819	1.00	21.83
ATOM	2167	CG	HIS	A	279	22.342	30.379	25.738	1.00	23.50
ATOM	2168	ND1	HIS	A	279	23.713	30.494	25.785	1.00	24.98
ATOM	2169	CD2	HIS	A	279	22.017	29.385	26.602	1.00	26.21
ATOM	2170	CE1	HIS	A	279	24.200	29.599	26.628	1.00	26.46
ATOM	2171	NE2	HIS	A	279	23.192	28.927	27.158	1.00	25.14
ATOM	2172	N	HIS	A	280	18.994	32.900	25.874	1.00	21.71
ATOM	2173	CA	HIS	A	280	17.736	32.956	26.602	1.00	21.78
ATOM	2174	C	HIS	A	280	16.890	31.860	25.951	1.00	21.70
ATOM	2175	O	HIS	A	280	16.722	31.846	24.736	1.00	21.40
ATOM	2176	CB	HIS	A	280	17.104	34.322	26.374	1.00	21.20
ATOM	2177	CG	HIS	A	280	15.642	34.367	26.626	1.00	21.64
ATOM	2178	ND1	HIS	A	280	15.100	35.028	27.705	1.00	22.67
ATOM	2179	CD2	HIS	A	280	14.600	33.826	25.951	1.00	23.33
ATOM	2180	CE1	HIS	A	280	13.786	34.891	27.684	1.00	23.20
ATOM	2181	NE2	HIS	A	280	13.457	34.158	26.635	1.00	23.30
ATOM	2182	N	ILE	A	281	16.368	30.934	26.733	1.00	22.98
ATOM	2183	CA	ILE	A	281	15.658	29.779	26.158	1.00	23.36
ATOM	2184	C	ILE	A	281	14.313	29.606	26.800	1.00	23.58
ATOM	2185	O	ILE	A	281	14.209	29.592	28.014	1.00	23.63
ATOM	2186	CB	ILE	A	281	16.512	28.525	26.301	1.00	23.87

[illegible]

ATOM	2187	CG1	ILE	A	281	17.771	28.675	25.435	1.00	25.77
ATOM	2188	CG2	ILE	A	281	15.748	27.261	25.863	1.00	23.54
ATOM	2189	CD1	ILE	A	281	18.752	27.659	25.690	1.00	29.36
ATOM	2190	N	GLU	A	282	13.268	29.510	25.982	1.00	23.92
ATOM	2191	CA	GLU	A	282	11.923	29.337	26.525	1.00	24.26
ATOM	2192	C	GLU	A	282	11.113	28.234	25.834	1.00	23.82
ATOM	2193	O	GLU	A	282	11.115	28.107	24.606	1.00	24.00
ATOM	2194	CB	GLU	A	282	11.166	30.681	26.490	1.00	24.35
ATOM	2195	CG	GLU	A	282	10.887	31.249	25.112	1.00	25.59
ATOM	2196	CD	GLU	A	282	10.320	32.668	25.168	1.00	25.45
ATOM	2197	OE1	GLU	A	282	10.861	33.494	25.929	1.00	26.04
ATOM	2198	OE2	GLU	A	282	9.332	32.954	24.451	1.00	25.08
ATOM	2199	N	SER	A	283	10.460	27.419	26.648	1.00	23.28
ATOM	2200	CA	SER	A	283	9.573	26.373	26.166	1.00	23.66
ATOM	2201	C	SER	A	283	8.257	27.054	25.787	1.00	23.86
ATOM	2202	O	SER	A	283	7.678	27.780	26.593	1.00	23.20
ATOM	2203	CB	SER	A	283	9.341	25.321	27.247	1.00	23.36
ATOM	2204	OG	SER	A	283	10.473	24.497	27.401	1.00	23.30
ATOM	2205	N	LEU	A	284	7.792	26.837	24.564	1.00	24.56
ATOM	2206	CA	LEU	A	284	6.626	27.583	24.084	1.00	25.62
ATOM	2207	C	LEU	A	284	5.401	27.487	25.004	1.00	25.52
ATOM	2208	O	LEU	A	284	5.114	26.431	25.589	1.00	24.71
ATOM	2209	CB	LEU	A	284	6.237	27.157	22.680	1.00	25.64
ATOM	2210	CG	LEU	A	284	7.274	27.308	21.576	1.00	27.12
ATOM	2211	CD1	LEU	A	284	6.591	27.546	20.242	1.00	28.93
ATOM	2212	CD2	LEU	A	284	8.248	28.364	21.851	1.00	27.08
ATOM	2213	N	LEU	A	285	4.713	28.615	25.144	1.00	25.85
ATOM	2214	CA	LEU	A	285	3.490	28.676	25.942	1.00	26.84
ATOM	2215	C	LEU	A	285	2.507	27.691	25.345	1.00	27.10
ATOM	2216	O	LEU	A	285	2.424	27.565	24.139	1.00	27.10
ATOM	2217	CB	LEU	A	285	2.875	30.071	25.906	1.00	26.48
ATOM	2218	CG	LEU	A	285	3.709	31.225	26.438	1.00	27.70
ATOM	2219	CD1	LEU	A	285	3.075	32.554	26.026	1.00	28.22
ATOM	2220	CD2	LEU	A	285	3.845	31.125	27.928	1.00	28.39
ATOM	2221	N	ASN	A	286	1.779	26.987	26.196	1.00	27.81
ATOM	2222	CA	ASN	A	286	0.790	26.007	25.753	1.00	28.27
ATOM	2223	C	ASN	A	286	1.313	24.977	24.744	1.00	27.42
ATOM	2224	O	ASN	A	286	0.555	24.481	23.922	1.00	26.50
ATOM	2225	CB	ASN	A	286	-0.417	26.741	25.160	1.00	29.08
ATOM	2226	CG	ASN	A	286	-0.931	27.846	26.078	1.00	32.15
ATOM	2227	OD1	ASN	A	286	-1.484	27.577	27.154	1.00	34.93
ATOM	2228	ND2	ASN	A	286	-0.725	29.099	25.670	1.00	35.91
ATOM	2229	N	GLY	A	287	2.608	24.672	24.799	1.00	26.54
ATOM	2230	CA	GLY	A	287	3.201	23.727	23.879	1.00	25.64
ATOM	2231	C	GLY	A	287	3.474	22.371	24.497	1.00	25.01
ATOM	2232	O	GLY	A	287	4.031	21.502	23.829	1.00	24.77
ATOM	2233	N	GLY	A	288	3.110	22.187	25.766	1.00	24.43
ATOM	2234	CA	GLY	A	288	3.357	20.922	26.455	1.00	23.99
ATOM	2235	C	GLY	A	288	4.804	20.844	26.935	1.00	24.11
ATOM	2236	O	GLY	A	288	5.546	21.810	26.786	1.00	22.82
ATOM	2237	N	ILE	A	289	5.211	19.695	27.472	1.00	23.99
ATOM	2238	CA	ILE	A	289	6.550	19.522	28.000	1.00	24.69
ATOM	2239	C	ILE	A	289	7.605	19.496	26.908	1.00	24.25
ATOM	2240	O	ILE	A	289	7.350	19.092	25.774	1.00	25.20
ATOM	2241	CB	ILE	A	289	6.691	18.234	28.836	1.00	25.39
ATOM	2242	CG1	ILE	A	289	6.702	17.006	27.941	1.00	27.43
ATOM	2243	CG2	ILE	A	289	5.613	18.151	29.921	1.00	26.30
ATOM	2244	CD1	ILE	A	289	7.255	15.752	28.650	1.00	30.57
ATOM	2245	N	THR	A	290	8.808	19.912	27.267	1.00	23.28
ATOM	2246	CA	THR	A	290	9.908	19.908	26.334	1.00	22.12
ATOM	2247	C	THR	A	290	11.008	18.984	26.824	1.00	21.52

[illegible]

ATOM	2309	C	TYR	A	297	33.506	21.660	25.554	1.00	24.51
ATOM	2310	O	TYR	A	297	33.188	22.825	25.323	1.00	23.73
ATOM	2311	CB	TYR	A	297	32.882	20.332	23.610	1.00	23.38
ATOM	2312	CG	TYR	A	297	32.284	19.092	23.021	1.00	23.42
ATOM	2313	CD1	TYR	A	297	30.940	19.053	22.641	1.00	23.90
ATOM	2314	CD2	TYR	A	297	33.056	17.971	22.808	1.00	21.43
ATOM	2315	CE1	TYR	A	297	30.394	17.920	22.089	1.00	23.32
ATOM	2316	CE2	TYR	A	297	32.515	16.826	22.250	1.00	22.05
ATOM	2317	CZ	TYR	A	297	31.186	16.806	21.905	1.00	21.88
ATOM	2318	OH	TYR	A	297	30.644	15.669	21.365	1.00	21.30
ATOM	2319	N	LYS	A	298	34.637	21.334	26.160	1.00	25.35
ATOM	2320	CA	LYS	A	298	35.589	22.362	26.536	1.00	26.48
ATOM	2321	C	LYS	A	298	36.070	23.017	25.268	1.00	26.23
ATOM	2322	O	LYS	A	298	36.224	22.364	24.249	1.00	24.85
ATOM	2323	CB	LYS	A	298	36.783	21.771	27.308	1.00	27.15
ATOM	2324	CG	LYS	A	298	36.474	21.471	28.777	1.00	30.00
ATOM	2325	CD	LYS	A	298	37.666	20.901	29.550	1.00	33.20
ATOM	2326	CE	LYS	A	298	37.232	20.493	30.969	1.00	36.09
ATOM	2327	NZ	LYS	A	298	38.299	19.766	31.764	1.00	39.03
ATOM	2328	N	GLY	A	299	36.321	24.313	25.329	1.00	27.13
ATOM	2329	CA	GLY	A	299	36.798	25.014	24.159	1.00	28.27
ATOM	2330	C	GLY	A	299	38.229	24.649	23.837	1.00	29.23
ATOM	2331	O	GLY	A	299	38.892	24.005	24.613	1.00	28.19
ATOM	2332	N	ALA	A	300	38.674	25.046	22.660	1.00	31.78
ATOM	2333	CA	ALA	A	300	40.046	24.852	22.233	1.00	34.32
ATOM	2334	C	ALA	A	300	40.992	25.610	23.155	1.00	36.37
ATOM	2335	O	ALA	A	300	40.573	26.430	23.956	1.00	35.96
ATOM	2336	CB	ALA	A	300	40.210	25.355	20.806	1.00	34.38
ATOM	2337	N	PRO	A	301	42.280	25.349	23.025	1.00	39.70
ATOM	2338	CA	PRO	A	301	43.278	26.051	23.845	1.00	41.64
ATOM	2339	C	PRO	A	301	43.465	27.482	23.386	1.00	43.35
ATOM	2340	O	PRO	A	301	43.220	27.803	22.228	1.00	43.86
ATOM	2341	CB	PRO	A	301	44.566	25.275	23.572	1.00	41.46
ATOM	2342	CG	PRO	A	301	44.382	24.736	22.199	1.00	41.25
ATOM	2343	CD	PRO	A	301	42.906	24.416	22.068	1.00	40.04
ATOM	2344	N	THR	A	302	43.900	28.328	24.300	1.00	45.97
ATOM	2345	CA	THR	A	302	44.214	29.715	24.001	1.00	47.77
ATOM	2346	C	THR	A	302	45.593	29.713	23.362	1.00	48.92
ATOM	2347	O	THR	A	302	46.532	29.251	24.006	1.00	49.23
ATOM	2348	CB	THR	A	302	44.282	30.469	25.322	1.00	48.10
ATOM	2349	OG1	THR	A	302	43.006	30.416	25.976	1.00	49.37
ATOM	2350	CG2	THR	A	302	44.520	31.924	25.100	1.00	48.91
ATOM	2351	N	PRO	A	303	45.761	30.248	22.148	1.00	50.22
ATOM	2352	CA	PRO	A	303	47.064	30.150	21.474	1.00	50.62
ATOM	2353	C	PRO	A	303	48.191	30.728	22.316	1.00	50.64
ATOM	2354	O	PRO	A	303	47.883	31.390	23.307	1.00	50.98
ATOM	2355	CB	PRO	A	303	46.877	30.986	20.202	1.00	50.93
ATOM	2356	CG	PRO	A	303	45.407	30.999	19.968	1.00	50.70
ATOM	2357	CD	PRO	A	303	44.814	31.064	21.360	1.00	50.45
ATOM	2358	N	GLU	A	307	46.978	37.074	18.830	1.00	53.59
ATOM	2359	CA	GLU	A	307	46.946	38.016	17.714	1.00	53.82
ATOM	2360	C	GLU	A	307	45.902	39.090	17.922	1.00	52.97
ATOM	2361	O	GLU	A	307	44.792	38.810	18.358	1.00	53.64
ATOM	2362	CB	GLU	A	307	46.672	37.308	16.371	1.00	54.37
ATOM	2363	CG	GLU	A	307	46.875	38.231	15.159	1.00	56.16
ATOM	2364	CD	GLU	A	307	46.966	37.504	13.822	1.00	57.96
ATOM	2365	OE1	GLU	A	307	46.621	36.303	13.738	1.00	59.31
ATOM	2366	OE2	GLU	A	307	47.388	38.150	12.839	1.00	59.69
ATOM	2367	N	TYR	A	308	46.267	40.324	17.610	1.00	51.95
ATOM	2368	CA	TYR	A	308	45.343	41.436	17.702	1.00	51.19
ATOM	2369	C	TYR	A	308	44.693	41.603	16.337	1.00	50.27

C
O
C
C
C
C
C
C
C
O
N
C
C
O
C
C
C
C
N
N
C
C
O
N
C
C
C
O
C
N
C
C
C
O
C
C
C
C
N
C
C
C
O
C
C
C
C
N
C
C
C
O
C
C
C
O
N
C
C

[illegible]

ATOM	2431	NZ	LYS	A	315	38.206	45.270	22.318	1.00	41.38
ATOM	2432	N	VAL	A	316	36.079	39.314	27.330	1.00	23.78
ATOM	2433	CA	VAL	A	316	35.258	38.796	28.396	1.00	23.67
ATOM	2434	C	VAL	A	316	36.138	37.937	29.329	1.00	23.61
ATOM	2435	O	VAL	A	316	36.094	38.075	30.558	1.00	22.63
ATOM	2436	CB	VAL	A	316	34.092	37.961	27.873	1.00	23.67
ATOM	2437	CG1	VAL	A	316	33.315	37.370	29.037	1.00	23.53
ATOM	2438	CG2	VAL	A	316	33.159	38.816	27.005	1.00	23.69
ATOM	2439	N	ALA	A	317	36.940	37.065	28.732	1.00	22.51
ATOM	2440	CA	ALA	A	317	37.834	36.207	29.507	1.00	22.28
ATOM	2441	C	ALA	A	317	38.730	37.067	30.391	1.00	21.32
ATOM	2442	O	ALA	A	317	38.926	36.783	31.556	1.00	21.45
ATOM	2443	CB	ALA	A	317	38.680	35.325	28.574	1.00	21.87
ATOM	2444	N	ILE	A	318	39.220	38.153	29.834	1.00	20.92
ATOM	2445	CA	ILE	A	318	40.091	39.053	30.574	1.00	21.36
ATOM	2446	C	ILE	A	318	39.375	39.662	31.784	1.00	21.54
ATOM	2447	O	ILE	A	318	39.930	39.697	32.895	1.00	20.69
ATOM	2448	CB	ILE	A	318	40.650	40.153	29.646	1.00	21.19
ATOM	2449	CG1	ILE	A	318	41.664	39.560	28.670	1.00	21.41
ATOM	2450	CG2	ILE	A	318	41.315	41.267	30.461	1.00	21.05
ATOM	2451	CD1	ILE	A	318	42.217	40.569	27.663	1.00	21.73
ATOM	2452	N	MET	A	319	38.146	40.122	31.569	1.00	21.31
ATOM	2453	CA	MET	A	319	37.400	40.774	32.632	1.00	21.99
ATOM	2454	C	MET	A	319	37.094	39.787	33.748	1.00	22.21
ATOM	2455	O	MET	A	319	37.219	40.114	34.937	1.00	23.25
ATOM	2456	CB	MET	A	319	36.133	41.446	32.108	1.00	21.66
ATOM	2457	CG	MET	A	319	36.369	42.703	31.233	1.00	21.49
ATOM	2458	SD	MET	A	319	34.787	43.525	30.787	1.00	23.05
ATOM	2459	CE	MET	A	319	34.192	42.395	29.453	1.00	22.63
ATOM	2460	N	ARG	A	320	36.736	38.568	33.379	1.00	21.99
ATOM	2461	CA	ARG	A	320	36.492	37.544	34.383	1.00	21.57
ATOM	2462	C	ARG	A	320	37.750	37.319	35.223	1.00	21.22
ATOM	2463	O	ARG	A	320	37.669	37.210	36.439	1.00	21.65
ATOM	2464	CB	ARG	A	320	36.083	36.237	33.726	1.00	20.79
ATOM	2465	CG	ARG	A	320	34.702	36.243	33.014	1.00	22.36
ATOM	2466	CD	ARG	A	320	34.300	34.857	32.452	1.00	21.24
ATOM	2467	NE	ARG	A	320	34.147	33.910	33.558	1.00	21.37
ATOM	2468	CZ	ARG	A	320	33.148	33.957	34.423	1.00	21.73
ATOM	2469	NH1	ARG	A	320	32.196	34.869	34.308	1.00	22.69
ATOM	2470	NH2	ARG	A	320	33.100	33.109	35.425	1.00	23.33
ATOM	2471	N	ASN	A	321	38.908	37.216	34.571	1.00	21.35
ATOM	2472	CA	ASN	A	321	40.159	36.947	35.282	1.00	21.20
ATOM	2473	C	ASN	A	321	40.510	38.078	36.239	1.00	21.02
ATOM	2474	O	ASN	A	321	40.958	37.828	37.364	1.00	21.80
ATOM	2475	CB	ASN	A	321	41.314	36.648	34.311	1.00	21.07
ATOM	2476	CG	ASN	A	321	41.263	35.209	33.752	1.00	22.81
ATOM	2477	OD1	ASN	A	321	40.821	34.306	34.431	1.00	23.88
ATOM	2478	ND2	ASN	A	321	41.713	35.018	32.521	1.00	20.59
ATOM	2479	N	ILE	A	322	40.336	39.320	35.802	1.00	20.31
ATOM	2480	CA	ILE	A	322	40.611	40.447	36.683	1.00	19.83
ATOM	2481	C	ILE	A	322	39.721	40.344	37.918	1.00	19.45
ATOM	2482	O	ILE	A	322	40.178	40.531	39.037	1.00	18.04
ATOM	2483	CB	ILE	A	322	40.327	41.775	35.980	1.00	20.01
ATOM	2484	CG1	ILE	A	322	41.320	42.009	34.849	1.00	21.62
ATOM	2485	CG2	ILE	A	322	40.351	42.930	36.989	1.00	20.64
ATOM	2486	CD1	ILE	A	322	42.782	41.987	35.278	1.00	23.99
ATOM	2487	N	GLU	A	323	38.444	40.035	37.716	1.00	18.87
ATOM	2488	CA	GLU	A	323	37.533	39.940	38.841	1.00	19.15
ATOM	2489	C	GLU	A	323	37.964	38.845	39.781	1.00	19.09
ATOM	2490	O	GLU	A	323	37.997	39.044	40.980	1.00	19.36
ATOM	2491	CB	GLU	A	323	36.095	39.721	38.366	1.00	19.19

N
N
C
C
O
C
C
C
N
C
C
O
C
N
C
C
O
C
C
C
N
C
C
C
O
C
C
C
N
C
C
C
O
C
C
C
N
C
C
C
O
C

ATOM	2492	CG	GLU	A	323	35.478	40.987	37.787	1.00	19.24	C
ATOM	2493	CD	GLU	A	323	34.096	40.764	37.189	1.00	17.51	C
ATOM	2494	OE1	GLU	A	323	33.140	40.470	37.922	1.00	17.44	O
ATOM	2495	OE2	GLU	A	323	33.982	40.901	35.974	1.00	16.60	O
ATOM	2496	N	LYS	A	324	38.357	37.711	39.227	1.00	19.47	N
ATOM	2497	CA	LYS	A	324	38.781	36.570	40.037	1.00	20.33	C
ATOM	2498	C	LYS	A	324	40.040	36.869	40.839	1.00	20.67	C
ATOM	2499	O	LYS	A	324	40.090	36.562	42.023	1.00	20.56	O
ATOM	2500	CB	LYS	A	324	39.040	35.334	39.172	1.00	20.11	C
ATOM	2501	CG	LYS	A	324	37.803	34.678	38.572	1.00	20.73	C
ATOM	2502	CD	LYS	A	324	38.238	33.413	37.761	1.00	21.68	C
ATOM	2503	CE	LYS	A	324	37.117	32.866	36.857	1.00	22.30	C
ATOM	2504	NZ	LYS	A	324	37.534	31.609	36.153	1.00	19.72	N
ATOM	2505	N	MET	A	325	41.045	37.456	40.194	1.00	21.37	N
ATOM	2506	CA	MET	A	325	42.305	37.782	40.860	1.00	22.78	C
ATOM	2507	C	MET	A	325	42.106	38.789	41.990	1.00	22.78	C
ATOM	2508	O	MET	A	325	42.734	38.694	43.047	1.00	22.48	O
ATOM	2509	CB	MET	A	325	43.308	38.369	39.865	1.00	23.14	C
ATOM	2510	CG	MET	A	325	43.964	37.359	38.978	1.00	27.30	C
ATOM	2511	SD	MET	A	325	44.699	38.147	37.511	1.00	35.72	S
ATOM	2512	CE	MET	A	325	45.804	39.221	38.338	1.00	36.16	C
ATOM	2513	N	LEU	A	326	41.248	39.765	41.752	1.00	23.21	N
ATOM	2514	CA	LEU	A	326	40.977	40.795	42.751	1.00	24.47	C
ATOM	2515	C	LEU	A	326	40.307	40.195	43.969	1.00	24.48	C
ATOM	2516	O	LEU	A	326	40.659	40.500	45.083	1.00	24.21	O
ATOM	2517	CB	LEU	A	326	40.087	41.863	42.159	1.00	24.63	C
ATOM	2518	CG	LEU	A	326	40.618	43.282	41.988	1.00	27.80	C
ATOM	2519	CD1	LEU	A	326	42.135	43.441	42.036	1.00	28.29	C
ATOM	2520	CD2	LEU	A	326	40.038	43.820	40.682	1.00	28.01	C
ATOM	2521	N	GLY	A	327	39.350	39.314	43.726	1.00	25.29	N
ATOM	2522	CA	GLY	A	327	38.663	38.608	44.782	1.00	25.73	C
ATOM	2523	C	GLY	A	327	39.644	37.854	45.638	1.00	25.86	C
ATOM	2524	O	GLY	A	327	39.597	37.931	46.863	1.00	25.83	O
ATOM	2525	N	GLU	A	328	40.556	37.137	45.002	1.00	26.02	N
ATOM	2526	CA	GLU	A	328	41.558	36.397	45.762	1.00	26.93	C
ATOM	2527	C	GLU	A	328	42.557	37.306	46.480	1.00	25.70	C
ATOM	2528	O	GLU	A	328	42.883	37.070	47.635	1.00	24.67	O
ATOM	2529	CB	GLU	A	328	42.293	35.423	44.853	1.00	27.76	C
ATOM	2530	CG	GLU	A	328	41.403	34.282	44.375	1.00	31.57	C
ATOM	2531	CD	GLU	A	328	41.635	32.992	45.146	1.00	37.39	C
ATOM	2532	OE1	GLU	A	328	42.048	33.058	46.335	1.00	41.34	O
ATOM	2533	OE2	GLU	A	328	41.430	31.902	44.552	1.00	42.69	O
ATOM	2534	N	ALA	A	329	43.024	38.356	45.816	1.00	25.20	N
ATOM	2535	CA	ALA	A	329	44.031	39.219	46.433	1.00	25.42	C
ATOM	2536	C	ALA	A	329	43.475	40.039	47.582	1.00	25.67	C
ATOM	2537	O	ALA	A	329	44.141	40.247	48.570	1.00	25.92	O
ATOM	2538	CB	ALA	A	329	44.670	40.109	45.409	1.00	25.31	C
ATOM	2539	N	LEU	A	330	42.241	40.496	47.467	1.00	26.51	N
ATOM	2540	CA	LEU	A	330	41.656	41.306	48.529	1.00	27.12	C
ATOM	2541	C	LEU	A	330	41.156	40.453	49.677	1.00	27.74	C
ATOM	2542	O	LEU	A	330	40.845	40.959	50.748	1.00	27.62	O
ATOM	2543	CB	LEU	A	330	40.517	42.138	47.984	1.00	27.19	C
ATOM	2544	CG	LEU	A	330	40.966	43.131	46.915	1.00	27.69	C
ATOM	2545	CD1	LEU	A	330	39.747	43.661	46.211	1.00	28.15	C
ATOM	2546	CD2	LEU	A	330	41.774	44.271	47.525	1.00	27.60	C
ATOM	2547	N	GLY	A	331	41.054	39.154	49.434	1.00	28.45	N
ATOM	2548	CA	GLY	A	331	40.637	38.222	50.457	1.00	29.04	C
ATOM	2549	C	GLY	A	331	39.154	38.239	50.748	1.00	29.16	C
ATOM	2550	O	GLY	A	331	38.698	37.505	51.615	1.00	29.67	O
ATOM	2551	N	ASN	A	332	38.409	39.088	50.052	1.00	29.16	N
ATOM	2552	CA	ASN	A	332	36.973	39.190	50.243	1.00	29.38	C

ATOM	2553	C	ASN	A	332	36.342	39.718	48.958	1.00	28.80	
ATOM	2554	O	ASN	A	332	36.579	40.847	48.570	1.00	28.58	C
ATOM	2555	CB	ASN	A	332	36.662	40.123	51.413	1.00	29.68	O
ATOM	2556	CG	ASN	A	332	35.181	40.214	51.702	1.00	31.38	C
ATOM	2557	OD1	ASN	A	332	34.368	39.630	50.991	1.00	35.54	C
ATOM	2558	ND2	ASN	A	332	34.820	40.944	52.751	1.00	33.23	O
ATOM	2559	N	PRO	A	333	35.504	38.922	48.325	1.00	28.45	N
ATOM	2560	CA	PRO	A	333	34.944	39.288	47.025	1.00	28.38	C
ATOM	2561	C	PRO	A	333	34.094	40.534	47.093	1.00	28.04	C
ATOM	2562	O	PRO	A	333	33.913	41.187	46.073	1.00	28.11	O
ATOM	2563	CB	PRO	A	333	34.071	38.093	46.640	1.00	28.23	C
ATOM	2564	CG	PRO	A	333	34.203	37.097	47.701	1.00	29.36	C
ATOM	2565	CD	PRO	A	333	34.999	37.641	48.822	1.00	28.91	C
ATOM	2566	N	GLN	A	334	33.556	40.844	48.262	1.00	27.75	N
ATOM	2567	CA	GLN	A	334	32.727	42.033	48.400	1.00	28.08	C
ATOM	2568	C	GLN	A	334	33.582	43.282	48.300	1.00	26.62	C
ATOM	2569	O	GLN	A	334	33.064	44.365	48.086	1.00	26.31	O
ATOM	2570	CB	GLN	A	334	31.857	41.989	49.686	1.00	28.89	C
ATOM	2571	CG	GLN	A	334	30.594	41.090	49.459	1.00	32.52	C
ATOM	2572	CD	GLN	A	334	29.523	41.114	50.556	1.00	36.22	C
ATOM	2573	OE1	GLN	A	334	29.500	42.004	51.421	1.00	39.82	O
ATOM	2574	NE2	GLN	A	334	28.612	40.127	50.503	1.00	38.24	N
ATOM	2575	N	GLU	A	335	34.894	43.138	48.414	1.00	25.23	N
ATOM	2576	CA	GLU	A	335	35.763	44.290	48.269	1.00	24.55	C
ATOM	2577	C	GLU	A	335	36.090	44.584	46.790	1.00	22.95	C
ATOM	2578	O	GLU	A	335	36.691	45.597	46.475	1.00	21.47	O
ATOM	2579	CB	GLU	A	335	37.038	44.091	49.088	1.00	25.61	C
ATOM	2580	CG	GLU	A	335	36.820	44.189	50.599	1.00	29.47	C
ATOM	2581	CD	GLU	A	335	38.115	44.280	51.369	1.00	34.49	C
ATOM	2582	OE1	GLU	A	335	38.964	45.124	51.004	1.00	39.50	O
ATOM	2583	OE2	GLU	A	335	38.288	43.521	52.350	1.00	40.05	O
ATOM	2584	N	VAL	A	336	35.654	43.722	45.877	1.00	21.52	N
ATOM	2585	CA	VAL	A	336	35.993	43.898	44.458	1.00	20.69	C
ATOM	2586	C	VAL	A	336	35.422	45.176	43.855	1.00	19.97	C
ATOM	2587	O	VAL	A	336	36.134	45.949	43.239	1.00	19.57	O
ATOM	2588	CB	VAL	A	336	35.594	42.683	43.653	1.00	20.59	C
ATOM	2589	CG1	VAL	A	336	35.746	42.933	42.179	1.00	21.20	C
ATOM	2590	CG2	VAL	A	336	36.467	41.513	44.067	1.00	21.28	C
ATOM	2591	N	GLY	A	337	34.146	45.421	44.080	1.00	19.82	N
ATOM	2592	CA	GLY	A	337	33.492	46.598	43.568	1.00	19.66	C
ATOM	2593	C	GLY	A	337	34.130	47.906	43.981	1.00	19.56	C
ATOM	2594	O	GLY	A	337	34.510	48.693	43.131	1.00	19.35	O
ATOM	2595	N	PRO	A	338	34.202	48.162	45.278	1.00	19.93	N
ATOM	2596	CA	PRO	A	338	34.846	49.383	45.790	1.00	19.88	C
ATOM	2597	C	PRO	A	338	36.272	49.631	45.254	1.00	19.22	C
ATOM	2598	O	PRO	A	338	36.591	50.758	44.908	1.00	18.43	O
ATOM	2599	CB	PRO	A	338	34.830	49.191	47.316	1.00	19.61	C
ATOM	2600	CG	PRO	A	338	33.625	48.380	47.555	1.00	20.33	C
ATOM	2601	CD	PRO	A	338	33.579	47.367	46.359	1.00	20.75	C
ATOM	2602	N	LEU	A	339	37.107	48.612	45.171	1.00	19.21	N
ATOM	2603	CA	LEU	A	339	38.416	48.814	44.566	1.00	19.65	C
ATOM	2604	C	LEU	A	339	38.283	49.219	43.081	1.00	19.35	C
ATOM	2605	O	LEU	A	339	38.927	50.188	42.621	1.00	17.91	O
ATOM	2606	CB	LEU	A	339	39.279	47.570	44.693	1.00	20.38	C
ATOM	2607	CG	LEU	A	339	40.745	47.814	44.291	1.00	21.96	C
ATOM	2608	CD1	LEU	A	339	41.681	47.151	45.220	1.00	25.12	C
ATOM	2609	CD2	LEU	A	339	40.991	47.293	42.899	1.00	22.54	C
ATOM	2610	N	LEU	A	340	37.420	48.512	42.345	1.00	19.18	N
ATOM	2611	CA	LEU	A	340	37.231	48.821	40.928	1.00	19.26	C
ATOM	2612	C	LEU	A	340	36.764	50.260	40.746	1.00	19.52	C
ATOM	2613	O	LEU	A	340	37.306	50.989	39.893	1.00	18.71	O

ATOM	2614	CB	LEU	A	340	36.260	47.857	40.273	1.00	19.85
ATOM	2615	CG	LEU	A	340	36.823	46.470	39.960	1.00	21.77
ATOM	2616	CD1	LEU	A	340	35.745	45.671	39.299	1.00	22.81
ATOM	2617	CD2	LEU	A	340	38.088	46.539	39.072	1.00	21.14
ATOM	2618	N	ASN	A	341	35.798	50.676	41.570	1.00	19.12
ATOM	2619	CA	ASN	A	341	35.296	52.052	41.536	1.00	20.43
ATOM	2620	C	ASN	A	341	36.367	53.086	41.865	1.00	19.90
ATOM	2621	O	ASN	A	341	36.474	54.110	41.206	1.00	19.81
ATOM	2622	CB	ASN	A	341	34.090	52.232	42.485	1.00	20.40
ATOM	2623	CG	ASN	A	341	32.814	51.659	41.898	1.00	24.06
ATOM	2624	OD1	ASN	A	341	32.390	52.073	40.829	1.00	31.36
ATOM	2625	ND2	ASN	A	341	32.229	50.666	42.561	1.00	25.82
ATOM	2626	N	THR	A	342	37.129	52.812	42.912	1.00	19.43
ATOM	2627	CA	THR	A	342	38.227	53.664	43.305	1.00	19.71
ATOM	2628	C	THR	A	342	39.230	53.781	42.154	1.00	19.45
ATOM	2629	O	THR	A	342	39.739	54.859	41.893	1.00	19.23
ATOM	2630	CB	THR	A	342	38.873	53.073	44.555	1.00	20.44
ATOM	2631	OG1	THR	A	342	38.030	53.322	45.700	1.00	21.13
ATOM	2632	CG2	THR	A	342	40.208	53.756	44.874	1.00	20.71
ATOM	2633	N	MET	A	343	39.464	52.691	41.427	1.00	19.49
ATOM	2634	CA	MET	A	343	40.381	52.723	40.287	1.00	19.90
ATOM	2635	C	MET	A	343	39.932	53.649	39.164	1.00	19.94
ATOM	2636	O	MET	A	343	40.775	54.344	38.567	1.00	18.43
ATOM	2637	CB	MET	A	343	40.543	51.347	39.664	1.00	19.92
ATOM	2638	CG	MET	A	343	41.701	50.556	40.115	1.00	23.07
ATOM	2639	SD	MET	A	343	42.163	49.194	38.959	1.00	24.93
ATOM	2640	CE	MET	A	343	41.013	48.220	39.315	1.00	27.48
ATOM	2641	N	ILE	A	344	38.629	53.643	38.837	1.00	20.84
ATOM	2642	CA	ILE	A	344	38.159	54.412	37.679	1.00	21.51
ATOM	2643	C	ILE	A	344	37.624	55.800	37.918	1.00	21.46
ATOM	2644	O	ILE	A	344	37.751	56.639	37.018	1.00	21.65
ATOM	2645	CB	ILE	A	344	37.089	53.641	36.805	1.00	22.58
ATOM	2646	CG1	ILE	A	344	35.714	53.723	37.435	1.00	24.65
ATOM	2647	CG2	ILE	A	344	37.506	52.220	36.555	1.00	23.63
ATOM	2648	CD1	ILE	A	344	34.635	53.407	36.511	1.00	28.35
ATOM	2649	N	LYS	A	345	37.009	56.092	39.062	1.00	21.48
ATOM	2650	CA	LYS	A	345	36.410	57.433	39.154	1.00	22.11
ATOM	2651	C	LYS	A	345	37.382	58.569	39.127	1.00	20.79
ATOM	2652	O	LYS	A	345	38.380	58.614	39.863	1.00	20.06
ATOM	2653	CB	LYS	A	345	35.405	57.656	40.279	1.00	23.49
ATOM	2654	CG	LYS	A	345	35.497	56.838	41.473	1.00	28.03
ATOM	2655	CD	LYS	A	345	34.120	56.183	41.683	1.00	30.07
ATOM	2656	CE	LYS	A	345	33.282	56.986	42.624	1.00	32.44
ATOM	2657	NZ	LYS	A	345	33.859	56.977	43.982	1.00	36.32
ATOM	2658	N	GLY	A	346	37.062	59.501	38.243	1.00	19.28
ATOM	2659	CA	GLY	A	346	37.911	60.636	38.013	1.00	18.60
ATOM	2660	C	GLY	A	346	39.222	60.254	37.350	1.00	17.41
ATOM	2661	O	GLY	A	346	40.090	61.079	37.229	1.00	18.04
ATOM	2662	N	ARG	A	347	39.353	59.024	36.898	1.00	17.62
ATOM	2663	CA	ARG	A	347	40.616	58.563	36.297	1.00	17.91
ATOM	2664	C	ARG	A	347	40.415	57.992	34.874	1.00	18.08
ATOM	2665	O	ARG	A	347	41.108	58.395	33.938	1.00	17.06
ATOM	2666	CB	ARG	A	347	41.284	57.532	37.213	1.00	17.51
ATOM	2667	CG	ARG	A	347	41.719	58.081	38.592	1.00	15.94
ATOM	2668	CD	ARG	A	347	43.179	57.649	38.985	1.00	19.49
ATOM	2669	NE	ARG	A	347	43.165	56.242	38.980	1.00	18.58
ATOM	2670	CZ	ARG	A	347	44.048	55.405	38.518	1.00	14.88
ATOM	2671	NH1	ARG	A	347	45.271	55.726	38.085	1.00	17.47
ATOM	2672	NH2	ARG	A	347	43.678	54.158	38.611	1.00	11.18
ATOM	2673	N	TYR	A	348	39.450	57.091	34.732	1.00	18.81
ATOM	2674	CA	TYR	A	348	39.098	56.480	33.444	1.00	20.50

C
C
C
C
N
C
C
O
C
C
O
N
N
C
C
O
C
O
C
C
S
C
N
C
C
O
C
C
C
C
C
N
N
C
C
O
N
C
C
O
C
C
C
N
N
N
C

ATOM	2675	C	TYR	A	348	37.637	56.691	33.036	1.00	22.07
ATOM	2676	O	TYR	A	348	37.233	56.236	31.978	1.00	22.11
ATOM	2677	CB	TYR	A	348	39.311	54.963	33.478	1.00	19.77
ATOM	2678	CG	TYR	A	348	40.753	54.545	33.561	1.00	20.36
ATOM	2679	CD1	TYR	A	348	41.540	54.483	32.422	1.00	19.91
ATOM	2680	CD2	TYR	A	348	41.335	54.225	34.780	1.00	18.20
ATOM	2681	CE1	TYR	A	348	42.859	54.099	32.489	1.00	19.81
ATOM	2682	CE2	TYR	A	348	42.662	53.841	34.863	1.00	19.47
ATOM	2683	CZ	TYR	A	348	43.425	53.785	33.711	1.00	19.43
ATOM	2684	OH	TYR	A	348	44.742	53.414	33.773	1.00	15.61
ATOM	2685	N	ASN	A	349	36.820	57.344	33.850	1.00	24.37
ATOM	2686	CA	ASN	A	349	35.414	57.472	33.456	1.00	26.66
ATOM	2687	C	ASN	A	349	34.941	58.854	33.092	1.00	28.03
ATOM	2688	O	ASN	A	349	35.663	59.847	32.992	1.00	27.79
ATOM	2689	CB	ASN	A	349	34.498	56.919	34.518	1.00	26.23
ATOM	2690	CG	ASN	A	349	34.463	57.764	35.748	1.00	27.89
ATOM	2691	OD1	ASN	A	349	35.247	58.718	35.927	1.00	27.38
ATOM	2692	ND2	ASN	A	349	33.545	57.409	36.645	1.00	32.46
ATOM	2693	OXT	ASN	A	349	33.736	58.963	32.900	1.00	32.44
TER	2694		ASN	A	349					
ATOM	2695	N	LEU	S	795	45.819	35.786	30.984	1.00	36.91
ATOM	2696	CA	LEU	S	795	44.711	36.756	31.250	1.00	37.26
ATOM	2697	C	LEU	S	795	43.553	36.418	30.323	1.00	37.49
ATOM	2698	O	LEU	S	795	42.391	36.451	30.712	1.00	36.90
ATOM	2699	CB	LEU	S	795	45.183	38.186	31.044	1.00	37.26
ATOM	2700	CG	LEU	S	795	44.683	39.204	32.074	1.00	37.32
ATOM	2701	CD1	LEU	S	795	44.775	38.671	33.479	1.00	37.34
ATOM	2702	CD2	LEU	S	795	45.479	40.493	32.006	1.00	37.95
ATOM	2703	N	THR	S	796	43.907	36.155	29.076	1.00	38.07
ATOM	2704	CA	THR	S	796	43.029	35.548	28.076	1.00	39.17
ATOM	2705	C	THR	S	796	42.608	34.088	28.347	1.00	38.93
ATOM	2706	O	THR	S	796	41.784	33.533	27.622	1.00	39.51
ATOM	2707	CB	THR	S	796	43.750	35.628	26.705	1.00	39.45
ATOM	2708	OG1	THR	S	796	43.287	34.597	25.850	1.00	41.87
ATOM	2709	CG2	THR	S	796	45.232	35.275	26.824	1.00	40.14
ATOM	2710	N	SER	S	797	43.154	33.451	29.369	1.00	38.98
ATOM	2711	CA	SER	S	797	42.770	32.067	29.654	1.00	39.24
ATOM	2712	C	SER	S	797	41.370	31.981	30.274	1.00	39.32
ATOM	2713	O	SER	S	797	40.901	32.905	30.939	1.00	38.28
ATOM	2714	CB	SER	S	797	43.775	31.398	30.574	1.00	38.98
ATOM	2715	OG	SER	S	797	43.613	31.884	31.891	1.00	40.40
ATOM	2716	N	TYR	S	798	40.713	30.851	30.059	1.00	39.93
ATOM	2717	CA	TYR	S	798	39.345	30.682	30.515	1.00	40.53
ATOM	2718	C	TYR	S	798	39.088	29.350	31.183	1.00	39.72
ATOM	2719	O	TYR	S	798	39.797	28.383	30.958	1.00	39.72
ATOM	2720	CB	TYR	S	798	38.377	30.881	29.351	1.00	41.28
ATOM	2721	CG	TYR	S	798	38.524	29.939	28.171	1.00	44.57
ATOM	2722	CD1	TYR	S	798	39.574	30.071	27.261	1.00	47.13
ATOM	2723	CD2	TYR	S	798	37.574	28.953	27.932	1.00	47.85
ATOM	2724	CE1	TYR	S	798	39.692	29.219	26.163	1.00	48.81
ATOM	2725	CE2	TYR	S	798	37.680	28.094	26.833	1.00	50.32
ATOM	2726	CZ	TYR	S	798	38.744	28.229	25.955	1.00	49.97
ATOM	2727	OH	TYR	S	798	38.835	27.387	24.866	1.00	49.58
ATOM	2728	N	ASP	S	799	38.051	29.306	32.005	1.00	39.16
ATOM	2729	CA	ASP	S	799	37.676	28.076	32.690	1.00	38.61
ATOM	2730	C	ASP	S	799	36.588	27.391	31.868	1.00	37.26
ATOM	2731	O	ASP	S	799	36.468	27.657	30.671	1.00	36.81
ATOM	2732	CB	ASP	S	799	37.186	28.399	34.105	1.00	39.29
ATOM	2733	CG	ASP	S	799	37.303	27.223	35.048	1.00	40.55
ATOM	2734	OD1	ASP	S	799	36.767	26.128	34.756	1.00	41.68
ATOM	2735	OD2	ASP	S	799	37.913	27.319	36.121	1.00	45.38

[illegible]

ATOM	2736	N	CYS	S	800	35.795	26.517	32.486	1.00	35.61
ATOM	2737	CA	CYS	S	800	34.781	25.787	31.732	1.00	34.66
ATOM	2738	C	CYS	S	800	33.439	25.714	32.450	1.00	34.03
ATOM	2739	O	CYS	S	800	32.749	24.708	32.388	1.00	33.81
ATOM	2740	CB	CYS	S	800	35.274	24.380	31.423	1.00	34.32
ATOM	2741	SG	CYS	S	800	35.538	23.392	32.911	1.00	33.88
ATOM	2742	N	GLU	S	801	33.051	26.792	33.101	1.00	33.69
ATOM	2743	CA	GLU	S	801	31.808	26.803	33.859	1.00	33.94
ATOM	2744	C	GLU	S	801	30.551	26.980	32.991	1.00	33.69
ATOM	2745	O	GLU	S	801	30.520	27.739	32.012	1.00	32.67
ATOM	2746	CB	GLU	S	801	31.886	27.877	34.942	1.00	34.03
ATOM	2747	CG	GLU	S	801	33.128	27.703	35.818	1.00	36.03
ATOM	2748	CD	GLU	S	801	33.095	28.557	37.065	1.00	36.15
ATOM	2749	OE1	GLU	S	801	32.408	28.163	38.013	1.00	35.62
ATOM	2750	OE2	GLU	S	801	33.751	29.619	37.090	1.00	38.69
ATOM	2751	N	VAL	S	802	29.519	26.242	33.377	1.00	33.95
ATOM	2752	CA	VAL	S	802	28.244	26.223	32.691	1.00	33.82
ATOM	2753	C	VAL	S	802	27.137	26.148	33.735	1.00	34.17
ATOM	2754	O	VAL	S	802	27.408	25.928	34.908	1.00	33.56
ATOM	2755	CB	VAL	S	802	28.139	24.972	31.781	1.00	33.63
ATOM	2756	CG1	VAL	S	802	29.263	24.948	30.753	1.00	32.61
ATOM	2757	CG2	VAL	S	802	28.165	23.682	32.624	1.00	33.77
ATOM	2758	N	ASN	S	803	25.887	26.304	33.300	1.00	35.05
ATOM	2759	CA	ASN	S	803	24.745	26.222	34.204	1.00	35.82
ATOM	2760	C	ASN	S	803	24.457	24.784	34.642	1.00	37.45
ATOM	2761	O	ASN	S	803	23.379	24.241	34.387	1.00	37.12
ATOM	2762	CB	ASN	S	803	23.490	26.836	33.573	1.00	35.36
ATOM	2763	CG	ASN	S	803	23.548	28.344	33.506	1.00	33.54
ATOM	2764	OD1	ASN	S	803	24.605	28.941	33.685	1.00	32.05
ATOM	2765	ND2	ASN	S	803	22.414	28.967	33.230	1.00	29.41
ATOM	2766	N	ALA	S	804	25.442	24.187	35.298	1.00	39.28
ATOM	2767	CA	ALA	S	804	25.340	22.846	35.855	1.00	41.13
ATOM	2768	C	ALA	S	804	26.566	22.620	36.734	1.00	42.43
ATOM	2769	O	ALA	S	804	27.638	23.153	36.464	1.00	42.47
ATOM	2770	CB	ALA	S	804	25.277	21.791	34.762	1.00	41.08
ATOM	2771	N	PRO	S	805	26.410	21.830	37.786	1.00	44.47
ATOM	2772	CA	PRO	S	805	27.527	21.527	38.697	1.00	45.68
ATOM	2773	C	PRO	S	805	28.568	20.620	38.053	1.00	46.82
ATOM	2774	O	PRO	S	805	28.273	19.918	37.092	1.00	47.40
ATOM	2775	CB	PRO	S	805	26.847	20.783	39.854	1.00	45.66
ATOM	2776	CG	PRO	S	805	25.588	20.206	39.261	1.00	45.01
ATOM	2777	CD	PRO	S	805	25.161	21.146	38.177	1.00	44.63
ATOM	2778	N	ILE	S	806	29.770	20.613	38.605	1.00	48.31
ATOM	2779	CA	ILE	S	806	30.847	19.766	38.098	1.00	49.34
ATOM	2780	C	ILE	S	806	30.661	18.323	38.555	1.00	49.57
ATOM	2781	O	ILE	S	806	29.992	18.063	39.560	1.00	50.13
ATOM	2782	CB	ILE	S	806	32.215	20.311	38.574	1.00	49.85
ATOM	2783	CG1	ILE	S	806	32.410	20.058	40.074	1.00	50.64
ATOM	2784	CG2	ILE	S	806	32.327	21.816	38.250	1.00	50.45
ATOM	2785	CD1	ILE	S	806	33.724	20.605	40.624	1.00	51.43
ATOM	2786	N	LEU	S	813	29.871	8.315	36.218	1.00	46.23
ATOM	2787	CA	LEU	S	813	30.588	8.788	35.034	1.00	46.32
ATOM	2788	C	LEU	S	813	29.685	9.635	34.140	1.00	45.87
ATOM	2789	O	LEU	S	813	28.463	9.511	34.185	1.00	45.43
ATOM	2790	CB	LEU	S	813	31.166	7.606	34.243	1.00	46.25
ATOM	2791	CG	LEU	S	813	32.057	6.663	35.059	1.00	46.61
ATOM	2792	CD1	LEU	S	813	32.444	5.423	34.230	1.00	46.51
ATOM	2793	CD2	LEU	S	813	33.292	7.406	35.579	1.00	45.98
ATOM	2794	N	GLN	S	814	30.309	10.503	33.351	1.00	45.81
ATOM	2795	CA	GLN	S	814	29.595	11.400	32.446	1.00	45.92
ATOM	2796	C	GLN	S	814	30.497	11.838	31.296	1.00	45.66

ATOM	2797	O	GLN	S	814	31.706	11.626	31.329	1.00	45.05	O
ATOM	2798	CB	GLN	S	814	29.123	12.643	33.198	1.00	46.09	C
ATOM	2799	CG	GLN	S	814	30.265	13.531	33.696	1.00	47.32	C
ATOM	2800	CD	GLN	S	814	29.781	14.722	34.520	1.00	49.22	C
ATOM	2801	OE1	GLN	S	814	29.165	14.541	35.569	1.00	50.40	O
ATOM	2802	NE2	GLN	S	814	30.070	15.937	34.053	1.00	49.45	N
ATOM	2803	N	GLY	S	815	29.898	12.476	30.294	1.00	45.88	N
ATOM	2804	CA	GLY	S	815	30.621	12.956	29.135	1.00	45.84	C
ATOM	2805	C	GLY	S	815	31.426	11.863	28.462	1.00	46.13	C
ATOM	2806	O	GLY	S	815	30.968	10.732	28.339	1.00	45.49	O
ATOM	2807	N	GLU	S	816	32.633	12.216	28.035	1.00	46.94	N
ATOM	2808	CA	GLU	S	816	33.548	11.285	27.382	1.00	48.03	C
ATOM	2809	C	GLU	S	816	33.721	9.994	28.186	1.00	48.62	C
ATOM	2810	O	GLU	S	816	33.823	8.910	27.614	1.00	48.54	O
ATOM	2811	CB	GLU	S	816	34.920	11.948	27.184	1.00	48.06	C
ATOM	2812	CG	GLU	S	816	35.783	11.312	26.102	1.00	48.93	C
ATOM	2813	CD	GLU	S	816	37.189	11.887	26.062	1.00	49.77	C
ATOM	2814	OE1	GLU	S	816	38.031	11.413	26.848	1.00	50.86	O
ATOM	2815	OE2	GLU	S	816	37.457	12.802	25.247	1.00	49.55	O
ATOM	2816	N	GLU	S	817	33.737	10.112	29.509	1.00	49.67	N
ATOM	2817	CA	GLU	S	817	33.947	8.957	30.373	1.00	50.75	C
ATOM	2818	C	GLU	S	817	32.786	7.982	30.309	1.00	51.10	C
ATOM	2819	O	GLU	S	817	32.989	6.771	30.278	1.00	51.12	O
ATOM	2820	CB	GLU	S	817	34.164	9.406	31.815	1.00	50.97	C
ATOM	2821	CG	GLU	S	817	35.476	10.140	32.032	1.00	52.22	C
ATOM	2822	CD	GLU	S	817	35.395	11.637	31.764	1.00	54.47	C
ATOM	2823	OE1	GLU	S	817	34.289	12.166	31.489	1.00	55.57	O
ATOM	2824	OE2	GLU	S	817	36.456	12.296	31.833	1.00	56.41	O
ATOM	2825	N	LEU	S	818	31.572	8.521	30.295	1.00	51.81	N
ATOM	2826	CA	LEU	S	818	30.376	7.706	30.204	1.00	52.37	C
ATOM	2827	C	LEU	S	818	30.399	6.929	28.901	1.00	53.14	C
ATOM	2828	O	LEU	S	818	30.216	5.714	28.894	1.00	53.00	O
ATOM	2829	CB	LEU	S	818	29.116	8.575	30.258	1.00	52.25	C
ATOM	2830	CG	LEU	S	818	27.786	7.814	30.176	1.00	52.17	C
ATOM	2831	CD1	LEU	S	818	27.661	6.836	31.335	1.00	52.37	C
ATOM	2832	CD2	LEU	S	818	26.596	8.749	30.173	1.00	51.16	C
ATOM	2833	N	LEU	S	819	30.660	7.643	27.809	1.00	54.00	N
ATOM	2834	CA	LEU	S	819	30.623	7.068	26.465	1.00	54.73	C
ATOM	2835	C	LEU	S	819	31.573	5.874	26.295	1.00	55.32	C
ATOM	2836	O	LEU	S	819	31.171	4.823	25.794	1.00	55.08	O
ATOM	2837	CB	LEU	S	819	30.929	8.162	25.426	1.00	54.66	C
ATOM	2838	CG	LEU	S	819	30.838	7.789	23.942	1.00	54.90	C
ATOM	2839	CD1	LEU	S	819	29.480	7.239	23.559	1.00	54.56	C
ATOM	2840	CD2	LEU	S	819	31.170	8.997	23.088	1.00	55.68	C
ATOM	2841	N	ARG	S	820	32.821	6.036	26.725	1.00	55.92	N
ATOM	2842	CA	ARG	S	820	33.817	4.987	26.576	1.00	56.75	C
ATOM	2843	C	ARG	S	820	33.488	3.762	27.428	1.00	56.92	C
ATOM	2844	O	ARG	S	820	33.578	2.617	26.951	1.00	56.86	O
ATOM	2845	CB	ARG	S	820	35.207	5.530	26.915	1.00	57.13	C
ATOM	2846	CG	ARG	S	820	35.638	6.596	25.927	1.00	58.55	C
ATOM	2847	CD	ARG	S	820	37.073	7.038	26.036	1.00	60.19	C
ATOM	2848	NE	ARG	S	820	37.336	8.126	25.096	1.00	62.23	N
ATOM	2849	CZ	ARG	S	820	38.465	8.831	25.039	1.00	64.08	C
ATOM	2850	NH1	ARG	S	820	39.475	8.567	25.865	1.00	64.81	N
ATOM	2851	NH2	ARG	S	820	38.586	9.805	24.142	1.00	64.58	N
ATOM	2852	N	ALA	S	821	33.109	4.003	28.682	1.00	56.90	N
ATOM	2853	CA	ALA	S	821	32.713	2.919	29.567	1.00	56.79	C
ATOM	2854	C	ALA	S	821	31.611	2.106	28.880	1.00	56.74	C
ATOM	2855	O	ALA	S	821	31.665	0.873	28.843	1.00	56.66	O
ATOM	2856	CB	ALA	S	821	32.236	3.463	30.899	1.00	56.66	C
ATOM	2857	N	LEU	S	822	30.629	2.806	28.317	1.00	56.51	N

ATOM	2858	CA	LEU S 822	29.537	2.155	27.607	1.00	56.58	
ATOM	2859	C	LEU S 822	30.031	1.473	26.338	1.00	56.66	C
ATOM	2860	O	LEU S 822	29.482	0.451	25.931	1.00	57.09	C
ATOM	2861	CB	LEU S 822	28.451	3.167	27.250	1.00	56.48	O
ATOM	2862	CG	LEU S 822	27.681	3.779	28.417	1.00	56.11	C
ATOM	2863	CD1	LEU S 822	26.543	4.612	27.870	1.00	55.96	C
ATOM	2864	CD2	LEU S 822	27.156	2.717	29.379	1.00	55.80	C
TER	2865		LEU S 822						C
HETATM	2866	FE	FE2 A1350	23.313	27.671	28.779	1.00	22.12	FE
HETATM	2867	C1	AKG A1351	22.355	25.315	27.747	1.00	25.61	C
HETATM	2868	O1	AKG A1351	23.449	25.880	27.756	1.00	27.58	O
HETATM	2869	O2	AKG A1351	22.172	24.103	27.197	1.00	27.99	O
HETATM	2870	C2	AKG A1351	21.128	25.999	28.365	1.00	24.14	C
HETATM	2871	O5	AKG A1351	21.211	27.117	28.854	1.00	23.66	O
HETATM	2872	C3	AKG A1351	19.829	25.231	28.280	1.00	23.46	C
HETATM	2873	C4	AKG A1351	18.717	25.967	29.008	1.00	22.15	C
HETATM	2874	C5	AKG A1351	17.351	25.435	28.649	1.00	23.90	C
HETATM	2875	O3	AKG A1351	17.136	24.674	27.706	1.00	23.27	O
HETATM	2876	O4	AKG A1351	16.353	25.844	29.406	1.00	25.34	O
HETATM	2877	S	SO4 A1352	0.196	25.255	43.681	1.00	83.69	S
HETATM	2878	O1	SO4 A1352	1.049	26.078	44.531	1.00	83.03	O
HETATM	2879	O2	SO4 A1352	1.028	24.391	42.840	1.00	82.84	O
HETATM	2880	O3	SO4 A1352	-0.643	24.431	44.542	1.00	83.90	O
HETATM	2881	O4	SO4 A1352	-0.630	26.114	42.830	1.00	82.82	O
HETATM	2882	S	SO4 A1353	1.937	28.607	29.759	1.00	80.69	S
HETATM	2883	O1	SO4 A1353	3.164	29.179	30.298	1.00	79.82	O
HETATM	2884	O2	SO4 A1353	2.228	27.552	28.793	1.00	79.89	O
HETATM	2885	O3	SO4 A1353	1.188	28.079	30.902	1.00	82.14	O
HETATM	2886	O4	SO4 A1353	1.145	29.630	29.081	1.00	81.49	O
HETATM	2887	O	HOH H 1	38.423	33.864	31.899	1.00	39.52	O
HETATM	2888	O	HOH H 2	38.025	25.366	29.554	1.00	64.59	O
HETATM	2889	O	HOH H 3	34.915	30.689	35.190	1.00	34.36	O
HETATM	2890	O	HOH H 4	20.482	27.037	33.306	1.00	55.20	O
HETATM	2891	O	HOH H 5	21.066	24.447	32.916	1.00	43.55	O
HETATM	2892	O	HOH H 6	29.978	24.394	35.721	1.00	43.81	O
HETATM	2893	O	HOH H 7	29.346	18.985	42.744	1.00	86.25	O
HETATM	2894	O	HOH H 8	35.530	13.904	24.157	1.00	42.65	O
HETATM	2895	O	HOH H 9	33.804	-1.383	26.877	1.00	65.05	O
HETATM	2896	O	HOH Z 1	11.560	21.626	13.846	1.00	41.47	O
HETATM	2897	O	HOH Z 2	9.590	21.877	12.314	1.00	61.59	O
HETATM	2898	O	HOH Z 3	1.321	21.339	7.657	1.00	58.53	O
HETATM	2899	O	HOH Z 4	3.579	13.365	8.778	1.00	47.77	O
HETATM	2900	O	HOH Z 5	4.515	16.855	3.766	1.00	51.50	O
HETATM	2901	O	HOH Z 6	2.462	19.552	5.161	1.00	56.40	O
HETATM	2902	O	HOH Z 7	1.251	29.413	13.184	1.00	52.18	O
HETATM	2903	O	HOH Z 8	2.053	32.304	13.875	1.00	71.43	O
HETATM	2904	O	HOH Z 9	11.574	44.907	14.867	1.00	67.18	O
HETATM	2905	O	HOH Z 10	11.615	3.238	17.221	1.00	63.99	O
HETATM	2906	O	HOH Z 11	3.752	32.951	32.375	1.00	72.66	O
HETATM	2907	O	HOH Z 12	4.803	37.611	27.421	1.00	63.47	O
HETATM	2908	O	HOH Z 13	11.007	35.734	30.393	1.00	34.95	O
HETATM	2909	O	HOH Z 14	15.551	46.392	24.481	1.00	43.01	O
HETATM	2910	O	HOH Z 15	12.231	41.979	15.720	1.00	53.94	O
HETATM	2911	O	HOH Z 16	13.868	4.815	17.661	1.00	48.46	O
HETATM	2912	O	HOH Z 17	15.860	30.606	12.755	1.00	44.45	O
HETATM	2913	O	HOH Z 18	13.462	22.030	7.390	1.00	59.18	O
HETATM	2914	O	HOH Z 19	14.706	26.336	13.845	1.00	51.42	O
HETATM	2915	O	HOH Z 20	17.028	29.994	7.603	1.00	64.07	O
HETATM	2916	O	HOH Z 21	21.135	23.988	3.773	1.00	46.32	O
HETATM	2917	O	HOH Z 22	27.581	31.130	6.026	1.00	64.13	O
HETATM	2918	O	HOH Z 23	27.341	22.242	43.414	1.00	79.43	O

HETATM	2919	O	HOH	Z	24	36.742	29.331	21.279	1.00	50.70	O
HETATM	2920	O	HOH	Z	25	30.029	33.533	9.206	1.00	50.33	O
HETATM	2921	O	HOH	Z	26	29.955	37.104	10.551	1.00	70.40	O
HETATM	2922	O	HOH	Z	27	18.215	15.129	13.036	1.00	37.33	O
HETATM	2923	O	HOH	Z	28	29.069	5.533	17.355	1.00	44.84	O
HETATM	2924	O	HOH	Z	29	18.941	14.771	16.383	1.00	31.41	O
HETATM	2925	O	HOH	Z	30	13.624	7.655	18.343	1.00	46.45	O
HETATM	2926	O	HOH	Z	31	5.649	12.667	27.758	1.00	44.15	O
HETATM	2927	O	HOH	Z	32	18.818	6.772	36.717	1.00	51.59	O
HETATM	2928	O	HOH	Z	33	7.620	14.589	19.463	1.00	50.93	O
HETATM	2929	O	HOH	Z	34	20.087	9.746	36.974	1.00	52.81	O
HETATM	2930	O	HOH	Z	35	21.912	13.173	44.511	1.00	59.64	O
HETATM	2931	O	HOH	Z	36	29.233	39.992	16.108	1.00	75.75	O
HETATM	2932	O	HOH	Z	37	33.785	44.067	25.671	1.00	62.06	O
HETATM	2933	O	HOH	Z	38	15.613	37.779	35.493	1.00	52.50	O
HETATM	2934	O	HOH	Z	39	8.070	38.292	35.056	1.00	60.61	O
HETATM	2935	O	HOH	Z	40	16.339	30.957	40.378	1.00	43.15	O
HETATM	2936	O	HOH	Z	41	28.116	27.147	37.617	1.00	59.20	O
HETATM	2937	O	HOH	Z	42	29.707	30.087	39.279	1.00	54.89	O
HETATM	2938	O	HOH	Z	43	28.116	24.509	42.048	1.00	63.13	O
HETATM	2939	O	HOH	Z	44	25.074	24.801	42.258	1.00	54.81	O
HETATM	2940	O	HOH	Z	45	33.873	31.493	39.077	1.00	45.97	O
HETATM	2941	O	HOH	Z	46	31.533	33.860	46.118	1.00	50.65	O
HETATM	2942	O	HOH	Z	47	13.319	35.957	31.390	1.00	44.72	O
HETATM	2943	O	HOH	Z	48	27.155	38.119	52.311	1.00	64.05	O
HETATM	2944	O	HOH	Z	49	24.587	38.767	49.612	1.00	50.58	O
HETATM	2945	O	HOH	Z	50	21.687	17.630	48.071	1.00	77.36	O
HETATM	2946	O	HOH	Z	51	21.437	14.872	40.880	1.00	60.20	O
HETATM	2947	O	HOH	Z	52	24.790	15.406	39.359	1.00	78.81	O
HETATM	2948	O	HOH	Z	53	23.347	17.356	36.625	1.00	52.48	O
HETATM	2949	O	HOH	Z	54	21.628	10.475	34.469	1.00	47.30	O
HETATM	2950	O	HOH	Z	55	18.013	-1.527	33.036	1.00	61.93	O
HETATM	2951	O	HOH	Z	56	16.101	-0.104	30.078	1.00	59.33	O
HETATM	2952	O	HOH	Z	57	26.268	5.539	16.988	1.00	38.42	O
HETATM	2953	O	HOH	Z	58	30.916	16.527	11.437	1.00	44.01	O
HETATM	2954	O	HOH	Z	59	32.683	13.953	20.664	1.00	50.04	O
HETATM	2955	O	HOH	Z	60	36.797	10.766	7.771	1.00	80.75	O
HETATM	2956	O	HOH	Z	61	33.878	26.222	17.133	1.00	45.23	O
HETATM	2957	O	HOH	Z	62	13.442	21.089	28.459	1.00	36.04	O
HETATM	2958	O	HOH	Z	63	3.999	21.370	30.471	1.00	47.33	O
HETATM	2959	O	HOH	Z	64	30.697	38.141	33.290	1.00	31.70	O
HETATM	2960	O	HOH	Z	65	26.005	26.456	26.227	1.00	28.76	O
HETATM	2961	O	HOH	Z	66	36.729	32.146	33.280	1.00	42.61	O
HETATM	2962	O	HOH	Z	67	35.846	25.574	27.896	1.00	34.95	O
HETATM	2963	O	HOH	Z	68	36.793	26.712	21.173	1.00	33.97	O
HETATM	2964	O	HOH	Z	69	17.427	17.022	18.148	1.00	31.28	O
HETATM	2965	O	HOH	Z	70	9.904	13.694	19.533	1.00	41.70	O
HETATM	2966	O	HOH	Z	71	5.361	16.931	22.051	1.00	43.04	O
HETATM	2967	O	HOH	Z	72	7.094	16.984	20.250	1.00	43.57	O
HETATM	2968	O	HOH	Z	73	6.562	22.961	22.902	1.00	42.74	O
HETATM	2969	O	HOH	Z	74	29.508	38.942	26.471	1.00	26.72	O
HETATM	2970	O	HOH	Z	75	30.732	39.209	19.135	1.00	37.64	O
HETATM	2971	O	HOH	Z	76	26.368	42.318	17.836	1.00	57.14	O
HETATM	2972	O	HOH	Z	77	27.688	44.616	31.257	1.00	30.61	O
HETATM	2973	O	HOH	Z	78	30.230	44.988	25.170	1.00	39.14	O
HETATM	2974	O	HOH	Z	79	27.780	48.720	30.030	1.00	38.89	O
HETATM	2975	O	HOH	Z	80	25.931	50.741	30.611	1.00	40.27	O
HETATM	2976	O	HOH	Z	81	18.521	38.529	36.775	1.00	43.87	O
HETATM	2977	O	HOH	Z	82	26.678	31.402	38.482	1.00	36.08	O
HETATM	2978	O	HOH	Z	83	30.586	30.409	36.592	1.00	32.57	O
HETATM	2979	O	HOH	Z	84	29.411	37.141	35.473	1.00	26.16	O

HETATM 2980	O	HOH Z 85	19.821	31.713	33.874	1.00	34.80	O
HETATM 2981	O	HOH Z 86	19.420	36.322	33.379	1.00	32.92	O
HETATM 2982	O	HOH Z 87	21.063	42.853	40.110	1.00	41.23	O
HETATM 2983	O	HOH Z 88	17.544	37.859	32.276	1.00	35.87	O
HETATM 2984	O	HOH Z 89	9.230	41.082	35.833	1.00	51.41	O
HETATM 2985	O	HOH Z 90	9.313	43.744	27.890	1.00	70.60	O
HETATM 2986	O	HOH Z 91	12.728	42.598	26.938	1.00	40.87	O
HETATM 2987	O	HOH Z 92	15.113	37.993	32.591	1.00	35.55	O
HETATM 2988	O	HOH Z 93	10.676	48.283	31.613	1.00	62.24	O
HETATM 2989	O	HOH Z 94	15.611	44.853	34.883	1.00	32.72	O
HETATM 2990	O	HOH Z 95	15.874	51.836	39.217	1.00	66.58	O
HETATM 2991	O	HOH Z 96	15.796	47.224	39.264	1.00	52.95	O
HETATM 2992	O	HOH Z 97	26.624	53.557	28.816	1.00	69.05	O
HETATM 2993	O	HOH Z 98	15.381	50.418	22.170	1.00	36.68	O
HETATM 2994	O	HOH Z 99	15.121	55.730	27.489	1.00	51.35	O
HETATM 2995	O	HOH Z 100	18.542	56.170	28.175	1.00	58.02	O
HETATM 2996	O	HOH Z 101	23.731	46.355	19.907	1.00	39.06	O
HETATM 2997	O	HOH Z 102	16.618	46.781	22.039	1.00	33.91	O
HETATM 2998	O	HOH Z 103	26.585	40.624	15.634	1.00	69.17	O
HETATM 2999	O	HOH Z 104	12.758	29.333	13.489	1.00	35.42	O
HETATM 3000	O	HOH Z 105	10.886	19.245	14.132	1.00	51.89	O
HETATM 3001	O	HOH Z 106	19.776	18.049	13.245	1.00	33.88	O
HETATM 3002	O	HOH Z 107	14.725	18.642	12.190	1.00	40.50	O
HETATM 3003	O	HOH Z 108	27.783	27.681	24.556	1.00	27.24	O
HETATM 3004	O	HOH Z 109	35.999	32.896	30.270	1.00	41.19	O
HETATM 3005	O	HOH Z 110	30.237	36.282	26.881	1.00	28.77	O
HETATM 3006	O	HOH Z 111	32.759	34.258	19.346	1.00	47.40	O
HETATM 3007	O	HOH Z 112	27.418	30.315	25.756	1.00	27.70	O
HETATM 3008	O	HOH Z 113	16.248	36.360	29.657	1.00	34.03	O
HETATM 3009	O	HOH Z 114	7.438	31.072	24.792	1.00	43.13	O
HETATM 3010	O	HOH Z 115	7.743	30.565	27.379	1.00	39.83	O
HETATM 3011	O	HOH Z 116	5.158	19.080	24.012	1.00	45.77	O
HETATM 3012	O	HOH Z 117	6.366	24.013	25.459	1.00	41.84	O
HETATM 3013	O	HOH Z 118	42.594	37.813	18.527	1.00	64.57	O
HETATM 3014	O	HOH Z 119	42.361	44.340	19.742	1.00	59.24	O
HETATM 3015	O	HOH Z 120	34.674	39.749	17.782	1.00	53.99	O
HETATM 3016	O	HOH Z 121	33.762	37.015	20.310	1.00	39.85	O
HETATM 3017	O	HOH Z 122	33.121	39.446	33.667	1.00	30.35	O
HETATM 3018	O	HOH Z 123	37.674	29.865	38.229	1.00	64.32	O
HETATM 3019	O	HOH Z 124	38.677	34.824	42.977	1.00	44.37	O
HETATM 3020	O	HOH Z 125	41.375	43.570	51.489	1.00	52.41	O
HETATM 3021	O	HOH Z 126	31.947	40.559	44.192	1.00	38.39	O
HETATM 3022	O	HOH Z 127	39.124	57.396	42.134	1.00	27.12	O
HETATM 3023	O	HOH Z 128	41.949	60.812	33.590	1.00	49.78	O
HETATM 3024	O	HOH Z 129	46.835	53.394	32.063	1.00	33.50	O
HETATM 3025	O	HOH Z 130	37.841	55.408	29.621	1.00	45.14	O
CONECT 1482	2866							
CONECT 1502	2866							
CONECT 2171	2866							
CONECT 2866	2871	2868	1482	2171	1502			
CONECT 2867	2868	2869	2870					
CONECT 2868	2867	2866						
CONECT 2869	2867							
CONECT 2870	2867	2871	2872					
CONECT 2871	2866	2870						
CONECT 2872	2870	2873						
CONECT 2873	2872	2874						
CONECT 2874	2873	2875	2876					
CONECT 2875	2874							
CONECT 2876	2874							
CONECT 2877	2878	2879	2880	2881				

CONECT 2878 2877
CONECT 2879 2877
CONECT 2880 2877
CONECT 2881 2877
CONECT 2882 2883 2884 2885 2886
CONECT 2883 2882
CONECT 2884 2882
CONECT 2885 2882
CONECT 2886 2882
MASTER 437 0 4 15 20 0 7 6 3023 2 24 31
END

Structure 3

Below are the coordinates for structure 3 (the 2.5 Å structure of FIH:Zn(II):NOG:CAD):

```

HEADER      TRANSCRIPTION ACTIVATOR/INHIBITOR          12-AUG-02   1H2M
TITLE       FACTOR INHIBITING HIF-1 ALPHA IN COMPLEX WITH HIF-1 ALPHA
TITLE       2 FRAGMENT PEPTIDE
COMPND      MOL_ID: 1;
COMPND      2 MOLECULE: FACTOR INHIBITING HIF1;
COMPND      3 SYNONYM: FIH1;
COMPND      4 CHAIN: A;
COMPND      5 ENGINEERED: YES;
COMPND      6 MOL_ID: 2;
COMPND      7 MOLECULE: HYPOXIA-INDUCIBLE FACTOR 1 ALPHA;
COMPND      8 SYNONYM: HIF-1 ALPHA, ARNT INTERACTING PROTEIN,
COMPND      9 MEMBER OF PAS PROTEIN 1, MOP1, HIF1 ALPHA, HIF1A.
COMPND     10 CHAIN: S;
COMPND     11 FRAGMENT: C-TERMINAL TRANSACTIVATION DOMAIN FRAGMENT
COMPND     12 RESIDUES 775 - 826
SOURCE      MOL_ID: 1;
SOURCE      2 ORGANISM_SCIENTIFIC: HOMO SAPIENS;
SOURCE      3 ORGANISM_COMMON: HUMAN;
SOURCE      4 EXPRESSION_SYSTEM: ESCHERICHIA COLI;
SOURCE      5 EXPRESSION_SYSTEM_STRAIN: BL21(DE3);
SOURCE      6 EXPRESSION_SYSTEM_PLASMID: PET28A(+);
SOURCE      7 MOL_ID: 2;
SOURCE      8 ORGANISM_SCIENTIFIC: HOMO SAPIENS;
SOURCE      9 ORGANISM_COMMON: HUMAN;
SOURCE     10 EXPRESSION_SYSTEM: ESCHERICHIA COLI;
SOURCE     11 EXPRESSION_SYSTEM_STRAIN: BL21(DE3);
SOURCE     12 EXPRESSION_SYSTEM_PLASMID: PGEX-GP-1
KEYWDS      FIH, HIF, DSBH, OXYGENASE, TRANSCRIPTION, HYPOXIA,
KEYWDS      2 2-OXOGLUTARATE, ASPARAGINYL HYDROXYLASE, HYDROXYLASE
EXPDTA      X-RAY DIFFRACTION
AUTHOR      J.M.ELKINS,K.S.HEWITSON,L.A.MCNEILL,I.SCHLEMMINGER,
AUTHOR      2 J.F.SEIBEL,C.J.SCHOFIELD
REVDAT      1 04-SEP-02 1H2M 0
JRNL        AUTH J.M.ELKINS,K.S.HEWITSON,L.A.MCNEILL,
JRNL        AUTH 2 I.SCHLEMMINGER,J.F.SEIBEL,C.J.SCHOFIELD
JRNL        TITL FIH:HIF-FRAGMENT COMPLEXES
JRNL        REF TO BE PUBLISHED
JRNL        REFN
REMARK      2
REMARK      2 RESOLUTION. 2.5  ANGSTROMS.
REMARK      3
REMARK      3 REFINEMENT.
REMARK      3 PROGRAM : REFMAC 5.0
REMARK      3 AUTHORS : MURSHUDOV,VAGIN,DODSON
REMARK      3
REMARK      3 REFINEMENT TARGET : MAXIMUM LIKELIHOOD
REMARK      3
REMARK      3 DATA USED IN REFINEMENT.
REMARK      3 RESOLUTION RANGE HIGH (ANGSTROMS) : 2.50
REMARK      3 RESOLUTION RANGE LOW (ANGSTROMS) : 18.00
REMARK      3 DATA CUTOFF (SIGMA(F)) : NONE
REMARK      3 COMPLETENESS FOR RANGE (%) : 99.68
REMARK      3 NUMBER OF REFLECTIONS : 18404
REMARK      3
REMARK      3 FIT TO DATA USED IN REFINEMENT.

```

```

REMARK 3 CROSS-VALIDATION METHOD : THROUGHOUT
REMARK 3 FREE R VALUE TEST SET SELECTION : RANDOM
REMARK 3 R VALUE (WORKING + TEST SET) : 0.19432
REMARK 3 R VALUE (WORKING SET) : 0.19185
REMARK 3 FREE R VALUE : 0.22491
REMARK 3 FREE R VALUE TEST SET SIZE (%) : 7.6
REMARK 3 FREE R VALUE TEST SET COUNT : 1516
REMARK 3
REMARK 3 FIT IN THE HIGHEST RESOLUTION BIN.
REMARK 3 TOTAL NUMBER OF BINS USED : 20
REMARK 3 BIN RESOLUTION RANGE HIGH : 2.500
REMARK 3 BIN RESOLUTION RANGE LOW : 2.564
REMARK 3 REFLECTION IN BIN (WORKING SET) : 1267
REMARK 3 BIN R VALUE (WORKING SET) : 0.227
REMARK 3 BIN FREE R VALUE SET COUNT : 106
REMARK 3 BIN FREE R VALUE : 0.297
REMARK 3
REMARK 3 NUMBER OF NON-HYDROGEN ATOMS USED IN REFINEMENT.
REMARK 3 ALL ATOMS : 2979
REMARK 3
REMARK 3 B VALUES.
REMARK 3 FROM WILSON PLOT (A**2) : NULL
REMARK 3 MEAN B VALUE (OVERALL, A**2) : 35.778
REMARK 3 OVERALL ANISOTROPIC B VALUE.
REMARK 3 B11 (A**2) : -0.68
REMARK 3 B22 (A**2) : -0.68
REMARK 3 B33 (A**2) : 1.35
REMARK 3 B12 (A**2) : 0.00
REMARK 3 B13 (A**2) : 0.00
REMARK 3 B23 (A**2) : 0.00
REMARK 3
REMARK 3 ESTIMATED OVERALL COORDINATE ERROR.
REMARK 3 ESU BASED ON R VALUE (A) : 0.334
REMARK 3 ESU BASED ON FREE R VALUE (A) : 0.233
REMARK 3 ESU BASED ON MAXIMUM LIKELIHOOD (A) : 0.224
REMARK 3 ESU FOR B VALUES BASED ON MAXIMUM LIKELIHOOD (A**2) : 9.825
REMARK 3
REMARK 3 CORRELATION COEFFICIENTS.
REMARK 3 CORRELATION COEFFICIENT FO-FC : 0.948
REMARK 3 CORRELATION COEFFICIENT FO-FC FREE : 0.935
REMARK 3
REMARK 3 RMS DEVIATIONS FROM IDEAL VALUES
REMARK 3 BOND LENGTHS REFINED ATOMS (A) : 2957 ; 0.017 ; 0.021
REMARK 3 BOND LENGTHS OTHERS (A) : 2546 ; 0.001 ; 0.020
REMARK 3 BOND ANGLES REFINED ATOMS (DEGREES) : 4022 ; 1.612 ; 1.948
REMARK 3 BOND ANGLES OTHERS (DEGREES) : 5944 ; 0.832 ; 3.000
REMARK 3 TORSION ANGLES, PERIOD 1 (DEGREES) : 350 ; 4.024 ; 3.000
REMARK 3 TORSION ANGLES, PERIOD 3 (DEGREES) : 512 ; 18.015 ; 15.000
REMARK 3 CHIRAL-CENTER RESTRAINTS (A**3) : 413 ; 0.097 ; 0.200
REMARK 3 GENERAL PLANES REFINED ATOMS (A) : 3315 ; 0.006 ; 0.020
REMARK 3 GENERAL PLANES OTHERS (A) : 602 ; 0.002 ; 0.020
REMARK 3 NON-BONDED CONTACTS REFINED ATOMS (A) : 731 ; 0.232 ; 0.300
REMARK 3 NON-BONDED CONTACTS OTHERS (A) : 2492 ; 0.214 ; 0.300
REMARK 3 H-BOND (X...Y) REFINED ATOMS (A) : 193 ; 0.173 ; 0.500
REMARK 3 H-BOND (X...Y) OTHERS (A) : 6 ; 0.126 ; 0.500
REMARK 3 POTENTIAL METAL-ION REFINED ATOMS (A) : 2 ; 0.054 ; 0.500
REMARK 3 SYMMETRY VDW REFINED ATOMS (A) : 15 ; 0.194 ; 0.300
REMARK 3 SYMMETRY VDW OTHERS (A) : 54 ; 0.255 ; 0.300
REMARK 3 SYMMETRY H-BOND REFINED ATOMS (A) : 7 ; 0.244 ; 0.500
REMARK 3 SYMMETRY H-BOND OTHERS (A) : 1 ; 0.053 ; 0.500
REMARK 3

```

REMARK 3 ISOTROPIC THERMAL FACTOR RESTRAINTS. COUNT RMS WEIGHT

REMARK 3 MAIN-CHAIN BOND REFINED ATOMS (A**2): 1767 ; 0.761 ; 1.500

REMARK 3 MAIN-CHAIN ANGLE REFINED ATOMS (A**2): 2846 ; 1.421 ; 2.000

REMARK 3 SIDE-CHAIN BOND REFINED ATOMS (A**2): 1190 ; 2.220 ; 3.000

REMARK 3 SIDE-CHAIN ANGLE REFINED ATOMS (A**2): 1176 ; 3.678 ; 4.500

REMARK 3 NCS RESTRAINTS STATISTICS

REMARK 3 NUMBER OF NCS GROUPS : NULL

REMARK 3 TLS DETAILS

REMARK 3 NUMBER OF TLS GROUPS : 1

REMARK 3 TLS GROUP : 1

REMARK 3 NUMBER OF COMPONENTS GROUP : 2

REMARK 3 COMPONENTS C SSSEQI TO C SSSEQI

REMARK 3 RESIDUE RANGE : A 15 A 451

REMARK 3 RESIDUE RANGE : S 795 S 822

REMARK 3 ORIGIN FOR THE GROUP (A): 22.5990 26.9200 28.6340

REMARK 3 T TENSOR

REMARK 3 T11: 0.1903 T22: 0.0302

REMARK 3 T33: 0.0452 T12: -0.0025

REMARK 3 T13: -0.0536 T23: 0.0309

REMARK 3 L TENSOR

REMARK 3 L11: 0.7638 L22: 2.2674

REMARK 3 L33: 1.0629 L12: 0.7977

REMARK 3 L13: 0.4200 L23: 1.0769

REMARK 3 S TENSOR

REMARK 3 S11: 0.0306 S12: -0.1225 S13: -0.0490

REMARK 3 S21: 0.1656 S22: 0.0303 S23: 0.0478

REMARK 3 S31: 0.2046 S32: 0.0231 S33: -0.0609

REMARK 3 BULK SOLVENT MODELLING.

REMARK 3 METHOD USED : BABINET MODEL WITH MASK

REMARK 3 PARAMETERS FOR MASK CALCULATION

REMARK 3 VDW PROBE RADIUS : 1.40

REMARK 3 ION PROBE RADIUS : 0.80

REMARK 3 SHRINKAGE RADIUS : 0.80

REMARK 3 OTHER REFINEMENT REMARKS:

REMARK 3 HYDROGENS HAVE BEEN ADDED IN THE RIDING POSITIONS

REMARK 4

REMARK 4 1H2M COMPLIES WITH FORMAT V. 2.3, 09-JULY-1998

REMARK 100

REMARK 100 THIS ENTRY HAS BEEN PROCESSED BY EBI ON 12-AUG-2002.

REMARK 100 THE EBI ID CODE IS EBI-11173.

REMARK 200

REMARK 200 EXPERIMENTAL DETAILS

REMARK 200 EXPERIMENT TYPE : X-RAY DIFFRACTION

REMARK 200 DATE OF DATA COLLECTION : 15-MAY-2002

REMARK 200 TEMPERATURE (KELVIN) : 100

REMARK 200 PH : 7.5

REMARK 200 NUMBER OF CRYSTALS USED : 1

REMARK 200

REMARK 200 SYNCHROTRON (Y/N) : Y

REMARK 200 RADIATION SOURCE : SRS BEAMLINE PX9.6

REMARK 200 BEAMLINE : PX9.6

REMARK 200 X-RAY GENERATOR MODEL : NULL

REMARK 200 MONOCHROMATIC OR LAUE (M/L) : M

REMARK 200 WAVELENGTH OR RANGE (A) : 0.87

REMARK 200 MONOCHROMATOR : NULL

REMARK 200 OPTICS : NULL

REMARK 200
 REMARK 200 DETECTOR TYPE : ADSC QUANTUM 4
 REMARK 200 DETECTOR MANUFACTURER : ADSC
 REMARK 200 INTENSITY-INTEGRATION SOFTWARE : MOSFLM
 REMARK 200 DATA SCALING SOFTWARE : SCALA
 REMARK 200
 REMARK 200 NUMBER OF UNIQUE REFLECTIONS : 20058
 REMARK 200 RESOLUTION RANGE HIGH (A) : 2.50
 REMARK 200 RESOLUTION RANGE LOW (A) : 87.71
 REMARK 200 REJECTION CRITERIA (SIGMA(I)) : NONE
 REMARK 200
 REMARK 200 OVERALL.
 REMARK 200 COMPLETENESS FOR RANGE (%) : 99.7
 REMARK 200 DATA REDUNDANCY : 6.5
 REMARK 200 R MERGE (I) : 0.050
 REMARK 200 R SYM (I) : NULL
 REMARK 200 <I/SIGMA(I)> FOR THE DATA SET : 10.7
 REMARK 200
 REMARK 200 IN THE HIGHEST RESOLUTION SHELL.
 REMARK 200 HIGHEST RESOLUTION SHELL, RANGE HIGH (A) : 2.50
 REMARK 200 HIGHEST RESOLUTION SHELL, RANGE LOW (A) : 2.64
 REMARK 200 COMPLETENESS FOR SHELL (%) : 97.9
 REMARK 200 DATA REDUNDANCY IN SHELL : 4.4
 REMARK 200 R MERGE FOR SHELL (I) : 0.289
 REMARK 200 R SYM FOR SHELL (I) : NULL
 REMARK 200 <I/SIGMA(I)> FOR SHELL : 2.6
 REMARK 200
 REMARK 200 DIFFRACTION PROTOCOL: SINGLE WAVELENGTH
 REMARK 200 METHOD USED TO DETERMINE THE STRUCTURE: MOLECULAR REPLACEMENT
 REMARK 200 SOFTWARE USED: NULL
 REMARK 200 STARTING MODEL: NULL
 REMARK 200
 REMARK 200 REMARK: NULL
 REMARK 280
 REMARK 280 CRYSTAL
 REMARK 280 SOLVENT CONTENT, VS (%): 63
 REMARK 280 MATTHEWS COEFFICIENT, VM (ANGSTROMS**3/DA): 3.4
 REMARK 280
 REMARK 280 CRYSTALLIZATION CONDITIONS: 1.2M AMMONIUM SULPHATE,
 REMARK 280 4% PEG400, 0.1M HEPES PH7.5, 11MG/ML PROTEIN WITH
 REMARK 280 1MM FE(II), 2.5MM NOG AND 2.5MM PEPTIDE
 REMARK 290
 REMARK 290 CRYSTALLOGRAPHIC SYMMETRY
 REMARK 290 SYMMETRY OPERATORS FOR SPACE GROUP: P 41 21 2
 REMARK 290
 REMARK 290 SYMOP SYMMETRY
 REMARK 290 NNNMMM OPERATOR
 REMARK 290 1555 X,Y,Z
 REMARK 290 2555 -X,-Y,1/2+Z
 REMARK 290 3555 1/2-Y,1/2+X,1/4+Z
 REMARK 290 4555 1/2+Y,1/2-X,3/4+Z
 REMARK 290 5555 1/2-X,1/2+Y,1/4-Z
 REMARK 290 6555 1/2+X,1/2-Y,3/4-Z
 REMARK 290 7555 Y,X,-Z
 REMARK 290 8555 -Y,-X,1/2-Z
 REMARK 290
 REMARK 290 WHERE NNN -> OPERATOR NUMBER
 REMARK 290 MMM -> TRANSLATION VECTOR
 REMARK 290
 REMARK 290 CRYSTALLOGRAPHIC SYMMETRY TRANSFORMATIONS
 REMARK 290 THE FOLLOWING TRANSFORMATIONS OPERATE ON THE ATOM/HETATM

REMARK 290 RECORDS IN THIS ENTRY TO PRODUCE CRYSTALLOGRAPHICALLY
 REMARK 290 RELATED MOLECULES.

REMARK 290	SMTRY1	1	1.000000	0.000000	0.000000	0.000000
REMARK 290	SMTRY2	1	0.000000	1.000000	0.000000	0.000000
REMARK 290	SMTRY3	1	0.000000	0.000000	1.000000	0.000000
REMARK 290	SMTRY1	2	-1.000000	0.000000	0.000000	0.000000
REMARK 290	SMTRY2	2	0.000000	-1.000000	0.000000	0.000000
REMARK 290	SMTRY3	2	0.000000	0.000000	1.000000	74.13000
REMARK 290	SMTRY1	3	0.000000	-1.000000	0.000000	43.12450
REMARK 290	SMTRY2	3	1.000000	0.000000	0.000000	43.12450
REMARK 290	SMTRY3	3	0.000000	0.000000	1.000000	37.06500
REMARK 290	SMTRY1	4	0.000000	1.000000	0.000000	43.12450
REMARK 290	SMTRY2	4	-1.000000	0.000000	0.000000	43.12450
REMARK 290	SMTRY3	4	0.000000	0.000000	1.000000	111.19500
REMARK 290	SMTRY1	5	-1.000000	0.000000	0.000000	43.12450
REMARK 290	SMTRY2	5	0.000000	1.000000	0.000000	43.12450
REMARK 290	SMTRY3	5	0.000000	0.000000	-1.000000	37.06500
REMARK 290	SMTRY1	6	1.000000	0.000000	0.000000	43.12450
REMARK 290	SMTRY2	6	0.000000	-1.000000	0.000000	43.12450
REMARK 290	SMTRY3	6	0.000000	0.000000	-1.000000	111.19500
REMARK 290	SMTRY1	7	0.000000	1.000000	0.000000	0.000000
REMARK 290	SMTRY2	7	1.000000	0.000000	0.000000	0.000000
REMARK 290	SMTRY3	7	0.000000	0.000000	-1.000000	0.000000
REMARK 290	SMTRY1	8	0.000000	-1.000000	0.000000	0.000000
REMARK 290	SMTRY2	8	-1.000000	0.000000	0.000000	0.000000
REMARK 290	SMTRY3	8	0.000000	0.000000	-1.000000	74.13000

REMARK 290

REMARK 290 REMARK: NULL

REMARK 300

REMARK 300 BIOMOLECULE: 1

REMARK 300 THIS ENTRY CONTAINS THE CRYSTALLOGRAPHIC ASYMMETRIC UNIT

REMARK 300 WHICH CONSISTS OF 2 CHAIN(S). SEE REMARK 350 FOR

REMARK 300 INFORMATION ON GENERATING THE BIOLOGICAL MOLECULE(S).

REMARK 300

REMARK 300 QUATERNARY STRUCTURE FOR THIS ENTRY: TETRAMERIC

REMARK 300

REMARK 300 THE PROTEIN IS A HOMODIMER FORMED BY CHAIN A.

REMARK 300 A HETERODIMERIC ASSOCIATION OF CHAIN A WITH CHAIN S

REMARK 300 PRODUCES A TETRAMER.

REMARK 300

REMARK 300 THE BURIED SURFACE AREA SHOWN BELOW IS AN AVERAGE

REMARK 300 CALCULATED FOR THE HETEROTETRAMER AND DOES NOT

REMARK 300 CORRESPOND TO THE BURIED SURFACE AREA FOR THE

REMARK 300 HOMODIMER OF CHAIN A

REMARK 300

REMARK 300 THE HETERO-ASSEMBLY DESCRIBED BY REMARK 350 APPEARS

REMARK 300 TO BE A CASE OF STRONG CRYSTAL PACKING WITH

REMARK 300 THE MEAN DIFFERENCE IN ACCESSIBLE SURFACE AREA PER

REMARK 300 CHAIN BETWEEN THE ISOLATED CHAIN AND THAT FOR

REMARK 300 THE CHAIN IN THE COMPLEX IS 2149.4 ANGSTROM**2

REMARK 350

REMARK 350 GENERATING THE BIOMOLECULE.

REMARK 350 COORDINATES FOR A COMPLETE MULTIMER REPRESENTING THE KNOWN

REMARK 350 BIOLOGICALLY SIGNIFICANT OLIGOMERIZATION STATE OF THE

REMARK 350 MOLECULE CAN BE GENERATED BY APPLYING BIOMT TRANSFORMATIONS

REMARK 350 GIVEN BELOW. BOTH NON-CRYSTALLOGRAPHIC AND

REMARK 350 CRYSTALLOGRAPHIC OPERATIONS ARE GIVEN.

REMARK 350

REMARK 350 BIOMOLECULE: 1

REMARK 350 APPLY THE FOLLOWING TO CHAINS: A, S

REMARK 350	BIOMT1	1	1.000000	0.000000	0.000000	0.000000
------------	--------	---	----------	----------	----------	----------

REMARK 350	BIOMT2	1	0.000000	1.000000	0.000000	0.000000
REMARK 350	BIOMT3	1	0.000000	0.000000	1.000000	0.000000
REMARK 350	BIOMT1	2	0.000000	-1.000000	0.000000	86.24900
REMARK 350	BIOMT2	2	-1.000000	0.000000	0.000000	86.24900
REMARK 350	BIOMT3	2	0.000000	0.000000	-1.000000	74.13000

REMARK 465

REMARK 465 MISSING RESIDUES

REMARK 465 THE FOLLOWING RESIDUES WERE NOT LOCATED IN THE

REMARK 465 EXPERIMENT. (M=MODEL NUMBER; RES=RESIDUE NAME; C=CHAIN

REMARK 465 IDENTIFIER; SSSEQ=SEQUENCE NUMBER; I=INSERTION CODE.)

REMARK 465

REMARK 465 M RES C SSSEQI

REMARK 465	MET	A	1
REMARK 465	ALA	A	2
REMARK 465	ALA	A	3
REMARK 465	THR	A	4
REMARK 465	ALA	A	5
REMARK 465	ALA	A	6
REMARK 465	GLU	A	7
REMARK 465	ALA	A	8
REMARK 465	VAL	A	9
REMARK 465	ALA	A	10
REMARK 465	SER	A	11
REMARK 465	GLY	A	12
REMARK 465	SER	A	13
REMARK 465	GLY	A	14
REMARK 465	LYS	A	304
REMARK 465	ARG	A	305
REMARK 465	ILE	A	306
REMARK 465	PRO	S	775
REMARK 465	SER	S	776
REMARK 465	ASP	S	777
REMARK 465	LEU	S	778
REMARK 465	ALA	S	779
REMARK 465	CYS	S	780
REMARK 465	ARG	S	781
REMARK 465	LEU	S	782
REMARK 465	LEU	S	783
REMARK 465	GLY	S	784
REMARK 465	GLN	S	785
REMARK 465	SER	S	786
REMARK 465	MET	S	787
REMARK 465	ASP	S	788
REMARK 465	GLU	S	789
REMARK 465	SER	S	790
REMARK 465	GLY	S	791
REMARK 465	LEU	S	792
REMARK 465	PRO	S	793
REMARK 465	GLN	S	794
REMARK 465	GLN	S	807
REMARK 465	GLY	S	808
REMARK 465	SER	S	809
REMARK 465	ARG	S	810
REMARK 465	ASN	S	811
REMARK 465	LEU	S	812
REMARK 465	ASP	S	823
REMARK 465	GLN	S	824
REMARK 465	VAL	S	825
REMARK 465	ASN	S	826

REMARK 470

REMARK 470 MISSING ATOM

PCT/GB2003/004492

REMARK 500 O ALA A 300 OH TYR S 798 2.09

REMARK 525

REMARK 525 SOLVENT

REMARK 525

REMARK 525 THE SOLVENT MOLECULES ARE GIVEN CHAIN IDENTIFIERS TO

REMARK 525 INDICATE THE PROTEIN CHAIN TO WHICH THEY ARE MOST CLOSELY

REMARK 525 ASSOCIATED WITH:

REMARK 525 PROTEIN CHAIN SOLVENT CHAIN

REMARK 525 A Z

REMARK 525 S H

REMARK 600

REMARK 600 HETEROGEN

REMARK 600

REMARK 600 FOR METAL ATOM ZN ZN A1350 THE COORDINATION ANGLES ARE:

REMARK 600 1 HIS 199A NE2

REMARK 600 2 ASP 201A OD2 103.4

REMARK 600 3 HIS 279A NE2 84.2 88.5

REMARK 600 4 OGA 1351A O2 169.0 87.2 99.0

REMARK 600 5 OGA 1351A O2' 86.3 169.1 97.3 82.8

REMARK 600 1 2 3 4

REMARK 700

REMARK 700 SHEET

REMARK 700 THE SHEET STRUCTURE OF THIS MOLECULE IS BIFURCATED. IN

REMARK 700 ORDER TO REPRESENT THIS FEATURE IN THE SHEET RECORDS BELOW,

REMARK 700 TWO SHEETS ARE DEFINED.

REMARK 800

REMARK 800 SITE

REMARK 800 SITE_IDENTIFIER: ZNA

REMARK 800 SITE_DESCRIPTION: ZN BINDING SITE FOR CHAIN A

REMARK 800

REMARK 800 SITE_IDENTIFIER: OGA

REMARK 800 SITE_DESCRIPTION: OGA BINDING SITE FOR CHAIN A

REMARK 800

REMARK 800 SITE_IDENTIFIER: SA1

REMARK 800 SITE_DESCRIPTION: SO4 BINDING SITE FOR CHAIN A

REMARK 800

REMARK 800 SITE_IDENTIFIER: SA2

REMARK 800 SITE_DESCRIPTION: SO4 BINDING SITE FOR CHAIN A

REMARK 900

REMARK 900 RELATED ENTRIES

REMARK 900 RELATED ID: 1D7G RELATED DB: PDB

REMARK 900 A MODEL FOR THE COMPLEX BETWEEN THE

REMARK 900 HYPOXIA-INDUCIBLE FACTOR-1 (HIF-1) AND ITS

REMARK 900 CONSENSUS DEOXYRIBONUCLEIC ACID SEQUENCE

REMARK 900 RELATED ID: 1H2K RELATED DB: PDB

REMARK 900 FACTOR INHIBITING HIF-1 ALPHA IN COMPLEX

REMARK 900 WITH HIF-1 ALPHA FRAGMENT PEPTIDE

REMARK 900 RELATED ID: 1H2L RELATED DB: PDB

REMARK 900 FACTOR INHIBITING HIF-1 ALPHA IN COMPLEX

REMARK 900 WITH HIF-1 ALPHA FRAGMENT PEPTIDE

REMARK 900 RELATED ID: 1H2N RELATED DB: PDB

REMARK 900 FACTOR INHIBITING HIF-1 ALPHA IN COMPLEX

REMARK 900 WITH HIF-1 ALPHA FRAGMENT PEPTIDE

REMARK 900 RELATED ID: 1L8C RELATED DB: PDB

REMARK 900 STRUCTURAL BASIS FOR HIF-1ALPHA/CBP

REMARK 900 RECOGNITION IN THECELLULAR HYPOXIC RESPONSE

REMARK 900 RELATED ID: 1LM8 RELATED DB: PDB

REMARK 900 STRUCTURE OF A HIF-1A-PVHL-ELONGINB-

REMARK 900 ELONGINC COMPLEX

REMARK 900 RELATED ID: 1LQB RELATED DB: PDB

REMARK 900 CRYSTAL STRUCTURE OF A HYDROXYLATED HIF-1

REMARK 900 ALPHA PEPTIDE BOUND TO THE PVHL/ELONGIN-C/
 REMARK 900 ELONGIN-B COMPLEX

DBREF 1H2M A 1 349 SWS Q969Q7 Q969Q7 1 349
 DBREF 1H2M S 775 826 SWS Q16665 HIFA_HUMAN 775 826

SEQRES 1 A 349 MET ALA ALA THR ALA ALA GLU ALA VAL ALA SER GLY SER
 SEQRES 2 A 349 GLY GLU PRO ARG GLU GLU ALA GLY ALA LEU GLY PRO ALA
 SEQRES 3 A 349 TRP ASP GLU SER GLN LEU ARG SER TYR SER PHE PRO THR
 SEQRES 4 A 349 ARG PRO ILE PRO ARG LEU SER GLN SER ASP PRO ARG ALA
 SEQRES 5 A 349 GLU GLU LEU ILE GLU ASN GLU GLU PRO VAL VAL LEU THR
 SEQRES 6 A 349 ASP THR ASN LEU VAL TYR PRO ALA LEU LYS TRP ASP LEU
 SEQRES 7 A 349 GLU TYR LEU GLN GLU ASN ILE GLY ASN GLY ASP PHE SER
 SEQRES 8 A 349 VAL TYR SER ALA SER THR HIS LYS PHE LEU TYR TYR ASP
 SEQRES 9 A 349 GLU LYS LYS MET ALA ASN PHE GLN ASN PHE LYS PRO ARG
 SEQRES 10 A 349 SER ASN ARG GLU GLU MET LYS PHE HIS GLU PHE VAL GLU
 SEQRES 11 A 349 LYS LEU GLN ASP ILE GLN GLN ARG GLY GLY GLU GLU ARG
 SEQRES 12 A 349 LEU TYR LEU GLN GLN THR LEU ASN ASP THR VAL GLY ARG
 SEQRES 13 A 349 LYS ILE VAL MET ASP PHE LEU GLY PHE ASN TRP ASN TRP
 SEQRES 14 A 349 ILE ASN LYS GLN GLN GLY LYS ARG GLY TRP GLY GLN LEU
 SEQRES 15 A 349 THR SER ASN LEU LEU LEU ILE GLY MET GLU GLY ASN VAL
 SEQRES 16 A 349 THR PRO ALA HIS TYR ASP GLU GLN GLN ASN PHE PHE ALA
 SEQRES 17 A 349 GLN ILE LYS GLY TYR LYS ARG CYS ILE LEU PHE PRO PRO
 SEQRES 18 A 349 ASP GLN PHE GLU CYS LEU TYR PRO TYR PRO VAL HIS HIS
 SEQRES 19 A 349 PRO CYS ASP ARG GLN SER GLN VAL ASP PHE ASP ASN PRO
 SEQRES 20 A 349 ASP TYR GLU ARG PHE PRO ASN PHE GLN ASN VAL VAL GLY
 SEQRES 21 A 349 TYR GLU THR VAL VAL GLY PRO GLY ASP VAL LEU TYR ILE
 SEQRES 22 A 349 PRO MET TYR TRP TRP HIS HIS ILE GLU SER LEU LEU ASN
 SEQRES 23 A 349 GLY GLY ILE THR ILE THR VAL ASN PHE TRP TYR LYS GLY
 SEQRES 24 A 349 ALA PRO THR PRO LYS ARG ILE GLU TYR PRO LEU LYS ALA
 SEQRES 25 A 349 HIS GLN LYS VAL ALA ILE MET ARG ASN ILE GLU LYS MET
 SEQRES 26 A 349 LEU GLY GLU ALA LEU GLY ASN PRO GLN GLU VAL GLY PRO
 SEQRES 27 A 349 LEU LEU ASN THR MET ILE LYS GLY ARG TYR ASN
 SEQRES 1 S 52 PRO SER ASP LEU ALA CYS ARG LEU LEU GLY GLN SER MET
 SEQRES 2 S 52 ASP GLU SER GLY LEU PRO GLN LEU THR SER TYR ASP CYS
 SEQRES 3 S 52 GLU VAL ASN ALA PRO ILE GLN GLY SER ARG ASN LEU LEU
 SEQRES 4 S 52 GLN GLY GLU GLU LEU LEU ARG ALA LEU ASP GLN VAL ASN

HET ZN A1350 1
 HET OGA A1351 10
 HET SO4 A1352 5
 HET SO4 A1353 5

HETNAM ZN ZINC ION
 HETNAM OGA N-OXALYOLGLYCINE
 HETNAM SO4 SULFATE ION

FORMUL 3 ZN ZN1 2+
 FORMUL 4 OGA C4 H5 N1 O5
 FORMUL 5 SO4 2(O4 S1 2-)
 FORMUL 6 HOH *99(H2 O1)

HELIX 1 1 ASP A 28 LEU A 32 5
 HELIX 2 2 ASP A 49 ASN A 58 1
 HELIX 3 3 VAL A 70 TRP A 76 5
 HELIX 4 4 ASP A 77 ILE A 85 1
 HELIX 5 5 ASP A 104 GLN A 112 5
 HELIX 6 6 LYS A 124 ARG A 138 1
 HELIX 7 7 GLY A 155 GLY A 164 1
 HELIX 8 8 ASN A 166 ARG A 177 1
 HELIX 9 9 PRO A 220 ASP A 222 5
 HELIX 10 10 GLN A 223 TYR A 228 1
 HELIX 11 11 PHE A 252 VAL A 258 5
 HELIX 12 12 LYS A 311 GLY A 331 1
 HELIX 13 13 ASN A 332 GLN A 334 5
 HELIX 14 14 GLU A 335 LYS A 345 1
 HELIX 15 15 GLN S 814 LEU S 822 1

SHEET	1	AA	5	THR	A	39		PRO	A	41	0								
SHEET	2	AA	5	GLY	A	260		VAL	A	265	1	O	GLY	A	260	N	ARG	A	40
SHEET	3	AA	5	LYS	A	214		PHE	A	219	-1	O	LYS	A	214	N	VAL	A	265
SHEET	4	AA	5	TRP	A	278		SER	A	283	-1	O	TRP	A	278	N	PHE	A	219
SHEET	5	AA	5	VAL	A	195		HIS	A	199	-1	O	THR	A	196	N	ILE	A	281
SHEET	1	AB	6	ARG	A	44		LEU	A	45	0								
SHEET	2	AB	6	VAL	A	62		LEU	A	64	1	O	VAL	A	63	N	LEU	A	45
SHEET	3	AB	6	VAL	A	270		ILE	A	273	-1	O	VAL	A	270	N	LEU	A	64
SHEET	4	AB	6	GLN	A	203		LYS	A	211	-1	O	ASN	A	205	N	ILE	A	273
SHEET	5	AB	6	THR	A	290		LYS	A	298	-1	O	ILE	A	291	N	ILE	A	210
SHEET	6	AB	6	LEU	A	182		SER	A	184	-1	N	THR	A	183	O	TRP	A	296
SHEET	1	AC	9	ARG	A	44		LEU	A	45	0								
SHEET	2	AC	9	VAL	A	62		LEU	A	64	1	O	VAL	A	63	N	LEU	A	45
SHEET	3	AC	9	VAL	A	270		ILE	A	273	-1	O	VAL	A	270	N	LEU	A	64
SHEET	4	AC	9	GLN	A	203		LYS	A	211	-1	O	ASN	A	205	N	ILE	A	273
SHEET	5	AC	9	THR	A	290		LYS	A	298	-1	O	ILE	A	291	N	ILE	A	210
SHEET	6	AC	9	LEU	A	186		GLY	A	190	-1	O	LEU	A	186	N	ASN	A	294
SHEET	7	AC	9	ARG	A	143		THR	A	149	-1	O	LEU	A	146	N	ILE	A	189
SHEET	8	AC	9	PHE	A	90		ALA	A	95	-1	O	SER	A	91	N	GLN	A	147
SHEET	9	AC	9	SER	A	118		MET	A	123	-1	O	ASN	A	119	N	SER	A	94
LINK		ZN											NE2	HIS	A	199		1555	1555
LINK		ZN											OD2	ASP	A	201		1555	1555
LINK		ZN											NE2	HIS	A	279		1555	1555
LINK		ZN											O2	OGA	A	1351		1555	1555
LINK		ZN											O2'	OGA	A	1351		1555	1555
CISPSP	1	TYR	A	308				PRO	A	309									
SITE	1	ZNA	3	HIS	A	199		ASP	A	201			HIS	A	279				
SITE	1	OGA	13	TYR	A	145		LEU	A	188			THR	A	196		HIS	A	199
SITE	2	OGA	13	ASP	A	201		ASN	A	205			PHE	A	207		LYS	A	214
SITE	3	OGA	13	HIS	A	279		ILE	A	281			ASN	A	294		TRP	A	296
SITE	4	OGA	13	HOH	Z	47													
SITE	1	SA1	4	ARG	A	138		GLY	A	140			GLU	A	141		GLU	A	142
SITE	1	SA2	5	ARG	A	143		GLU	A	192			GLY	A	193		LEU	A	285
SITE	2	SA2	5	ASN	A	286													
CRYST1	86.249			86.249															

ATOM	21	CZ	ARG	A	17	11.401	24.776	12.744	1.00	84.27
ATOM	22	NH1	ARG	A	17	11.980	25.968	12.584	1.00	83.33
ATOM	23	NH2	ARG	A	17	11.311	24.268	13.967	1.00	85.10
ATOM	24	N	GLU	A	18	5.796	24.209	8.421	1.00	74.01
ATOM	25	CA	GLU	A	18	4.692	23.274	8.319	1.00	72.99
ATOM	26	C	GLU	A	18	5.193	21.855	8.607	1.00	71.43
ATOM	27	O	GLU	A	18	6.210	21.421	8.061	1.00	70.71
ATOM	28	CB	GLU	A	18	4.033	23.352	6.938	1.00	73.29
ATOM	29	CG	GLU	A	18	3.333	24.677	6.659	1.00	73.91
ATOM	30	CD	GLU	A	18	1.949	24.794	7.283	1.00	74.30
ATOM	31	OE1	GLU	A	18	1.355	23.776	7.692	1.00	74.07
ATOM	32	OE2	GLU	A	18	1.444	25.929	7.356	1.00	75.76
ATOM	33	N	GLU	A	19	4.482	21.157	9.491	1.00	69.89
ATOM	34	CA	GLU	A	19	4.786	19.764	9.818	1.00	68.89
ATOM	35	C	GLU	A	19	4.393	18.854	8.661	1.00	66.65
ATOM	36	O	GLU	A	19	3.324	19.011	8.065	1.00	65.92
ATOM	37	CB	GLU	A	19	4.058	19.311	11.096	1.00	69.41
ATOM	38	CG	GLU	A	19	4.544	19.996	12.370	1.00	72.06
ATOM	39	CD	GLU	A	19	4.308	19.194	13.657	1.00	76.41
ATOM	40	OE1	GLU	A	19	3.855	18.016	13.609	1.00	78.21
ATOM	41	OE2	GLU	A	19	4.590	19.758	14.746	1.00	79.64
ATOM	42	N	ALA	A	20	5.283	17.919	8.349	1.00	64.54
ATOM	43	CA	ALA	A	20	5.067	16.921	7.303	1.00	63.03
ATOM	44	C	ALA	A	20	3.735	16.225	7.449	1.00	61.23
ATOM	45	O	ALA	A	20	3.303	15.903	8.556	1.00	60.61
ATOM	46	CB	ALA	A	20	6.177	15.889	7.312	1.00	62.82
ATOM	47	N	GLY	A	21	3.087	16.002	6.316	1.00	59.29
ATOM	48	CA	GLY	A	21	1.833	15.289	6.304	1.00	58.26
ATOM	49	C	GLY	A	21	0.651	16.200	6.530	1.00	57.44
ATOM	50	O	GLY	A	21	-0.416	15.739	6.873	1.00	56.42
ATOM	51	N	ALA	A	22	0.858	17.500	6.341	1.00	57.28
ATOM	52	CA	ALA	A	22	-0.182	18.509	6.492	1.00	56.92
ATOM	53	C	ALA	A	22	-0.737	18.544	7.909	1.00	56.53
ATOM	54	O	ALA	A	22	-1.926	18.737	8.108	1.00	56.32
ATOM	55	CB	ALA	A	22	-1.285	18.277	5.488	1.00	57.10
ATOM	56	N	LEU	A	23	0.133	18.369	8.898	1.00	56.25
ATOM	57	CA	LEU	A	23	-0.306	18.397	10.289	1.00	56.07
ATOM	58	C	LEU	A	23	-0.385	19.811	10.827	1.00	55.03
ATOM	59	O	LEU	A	23	-0.638	20.037	11.998	1.00	54.89
ATOM	60	CB	LEU	A	23	0.583	17.519	11.149	1.00	56.05
ATOM	61	CG	LEU	A	23	0.445	16.067	10.678	1.00	58.06
ATOM	62	CD1	LEU	A	23	1.307	15.080	11.478	1.00	59.22
ATOM	63	CD2	LEU	A	23	-1.030	15.653	10.735	1.00	59.53
ATOM	64	N	GLY	A	24	-0.202	20.768	9.941	1.00	54.17

ATOM	82	C	TRP	A	27	3.003	27.818	16.107	1.00	44.47
ATOM	83	O	TRP	A	27	2.985	27.981	14.901	1.00	44.30
ATOM	84	CB	TRP	A	27	4.719	26.727	17.568	1.00	44.91
ATOM	85	CG	TRP	A	27	5.916	26.910	16.737	1.00	42.67
ATOM	86	CD1	TRP	A	27	6.702	25.934	16.197	1.00	43.38
ATOM	87	CD2	TRP	A	27	6.411	28.129	16.255	1.00	38.70
ATOM	88	NE1	TRP	A	27	7.690	26.494	15.425	1.00	41.56
ATOM	89	CE2	TRP	A	27	7.532	27.847	15.452	1.00	39.59
ATOM	90	CE3	TRP	A	27	6.039	29.442	16.431	1.00	38.28
ATOM	91	CZ2	TRP	A	27	8.284	28.827	14.861	1.00	39.74
ATOM	92	CZ3	TRP	A	27	6.781	30.409	15.849	1.00	39.48
ATOM	93	CH2	TRP	A	27	7.884	30.104	15.057	1.00	40.17
ATOM	94	N	ASP	A	28	2.580	28.760	16.941	1.00	43.91
ATOM	95	CA	ASP	A	28	2.230	30.079	16.417	1.00	43.87
ATOM	96	C	ASP	A	28	2.750	31.161	17.305	1.00	42.33
ATOM	97	O	ASP	A	28	3.191	30.901	18.409	1.00	42.11
ATOM	98	CB	ASP	A	28	0.722	30.251	16.199	1.00	44.82
ATOM	99	CG	ASP	A	28	-0.052	30.162	17.462	1.00	47.87
ATOM	100	OD1	ASP	A	28	-0.240	31.236	18.108	1.00	50.91
ATOM	101	OD2	ASP	A	28	-0.501	29.054	17.877	1.00	49.06
ATOM	102	N	GLU	A	29	2.713	32.385	16.790	1.00	41.29
ATOM	103	CA	GLU	A	29	3.208	33.586	17.499	1.00	39.69
ATOM	104	C	GLU	A	29	2.685	33.711	18.917	1.00	37.94
ATOM	105	O	GLU	A	29	3.415	34.069	19.801	1.00	37.26
ATOM	106	CB	GLU	A	29	2.840	34.812	16.735	1.00	39.59
ATOM	107	N	SER	A	30	1.427	33.371	19.135	1.00	36.78
ATOM	108	CA	SER	A	30	0.810	33.558	20.443	1.00	36.13
ATOM	109	C	SER	A	30	1.501	32.756	21.517	1.00	35.30
ATOM	110	O	SER	A	30	1.252	32.968	22.678	1.00	35.21
ATOM	111	CB	SER	A	30	-0.686	33.180	20.406	1.00	35.84
ATOM	112	OG	SER	A	30	-0.901	31.762	20.450	1.00	36.93
ATOM	113	N	GLN	A	31	2.326	31.795	21.116	1.00	35.24
ATOM	114	CA	GLN	A	31	3.021	30.918	22.060	1.00	34.89
ATOM	115	C	GLN	A	31	4.366	31.506	22.487	1.00	34.60
ATOM	116	O	GLN	A	31	5.010	30.969	23.365	1.00	34.18
ATOM	117	CB	GLN	A	31	3.224	29.513	21.473	1.00	34.65
ATOM	118	CG	GLN	A	31	1.969	28.649	21.402	1.00	34.74
ATOM	119	CD	GLN	A	31	2.212	27.322	20.707	1.00	33.47
ATOM	120	OE1	GLN	A	31	2.215	27.249	19.476	1.00	32.57
ATOM	121	NE2	GLN	A	31	2.442	26.278	21.492	1.00	32.95
ATOM	122	N	LEU	A	32	4.753	32.632	21.895	1.00	34.85
ATOM	123	CA	LEU	A	32	6.016	33.293	22.212	1.00	35.01
ATOM	124	C	LEU	A	32	5.798	34.391	23.223	1.00	34.91
ATOM	125	O	LEU	A	32	4.834	35.134	23.125	1.00	35.83
ATOM	126	CB	LEU	A	32	6.631	33.885	20.945	1.00	34.96
ATOM	127	CG	LEU	A	32	6.995	32.860	19.849	1.00	36.03
ATOM	128	CD1	LEU	A	32	7.691	33.525	18.701	1.00	36.15
ATOM	129	CD2	LEU	A	32	7.855	31.780	20.377	1.00	35.18
ATOM	130	N	ARG	A	33	6.675	34.500	24.209	1.00	34.49
ATOM	131	CA	ARG	A	33	6.564	35.591	25.170	1.00	34.33
ATOM	132	C	ARG	A	33	7.005	36.867	24.460	1.00	34.37
ATOM	133	O	ARG	A	33	7.733	36.815	23.498	1.00	34.87
ATOM	134	CB	ARG	A	33	7.442	35.338	26.394	1.00	34.16
ATOM	135	CG	ARG	A	33	7.056	34.124	27.212	1.00	32.83
ATOM	136	CD	ARG	A	33	7.951	33.894	28.428	1.00	33.34
ATOM	137	NE	ARG	A	33	7.413	32.820	29.252	1.00	34.91
ATOM	138	CZ	ARG	A	33	6.445	32.963	30.137	1.00	36.53
ATOM	139	NH1	ARG	A	33	5.900	34.144	30.382	1.00	34.89
ATOM	140	NH2	ARG	A	33	6.027	31.905	30.795	1.00	39.39
ATOM	141	N	SER	A	34	6.608	38.022	24.947	1.00	34.36
ATOM	142	CA	SER	A	34	6.944	39.247	24.244	1.00	34.69

ATOM	143	C	SER A	34	8.002	40.055	24.987	1.00	33.26
ATOM	144	O	SER A	34	7.958	40.142	26.200	1.00	33.50
ATOM	145	CB	SER A	34	5.698	40.096	24.143	1.00	35.23
ATOM	146	OG	SER A	34	5.586	40.744	25.393	1.00	39.99
ATOM	147	N	TYR A	35	8.911	40.682	24.252	1.00	32.38
ATOM	148	CA	TYR A	35	10.080	41.329	24.843	1.00	32.03
ATOM	149	C	TYR A	35	10.339	42.675	24.193	1.00	32.52
ATOM	150	O	TYR A	35	9.763	42.979	23.172	1.00	32.78
ATOM	151	CB	TYR A	35	11.290	40.422	24.686	1.00	31.20
ATOM	152	CG	TYR A	35	11.139	39.161	25.482	1.00	30.69
ATOM	153	CD1	TYR A	35	10.935	39.215	26.853	1.00	30.40
ATOM	154	CD2	TYR A	35	11.186	37.915	24.875	1.00	30.34
ATOM	155	CE1	TYR A	35	10.804	38.054	27.609	1.00	31.74
ATOM	156	CE2	TYR A	35	11.050	36.741	25.617	1.00	32.05
ATOM	157	CZ	TYR A	35	10.858	36.812	26.991	1.00	31.98
ATOM	158	OH	TYR A	35	10.687	35.658	27.742	1.00	30.13
ATOM	159	N	SER A	36	11.240	43.463	24.761	1.00	33.06
ATOM	160	CA	SER A	36	11.469	44.838	24.289	1.00	32.95
ATOM	161	C	SER A	36	12.469	44.977	23.159	1.00	32.65
ATOM	162	O	SER A	36	12.705	46.084	22.674	1.00	33.04
ATOM	163	CB	SER A	36	11.979	45.698	25.438	1.00	32.92
ATOM	164	OG	SER A	36	13.272	45.325	25.844	1.00	33.05
ATOM	165	N	PHE A	37	13.041	43.880	22.711	1.00	31.44
ATOM	166	CA	PHE A	37	14.100	43.988	21.740	1.00	31.55
ATOM	167	C	PHE A	37	13.784	43.269	20.444	1.00	31.69
ATOM	168	O	PHE A	37	13.020	42.326	20.386	1.00	32.65
ATOM	169	CB	PHE A	37	15.421	43.431	22.339	1.00	30.68
ATOM	170	CG	PHE A	37	15.275	42.054	22.890	1.00	30.10
ATOM	171	CD1	PHE A	37	15.329	40.967	22.060	1.00	28.43
ATOM	172	CD2	PHE A	37	15.022	41.852	24.231	1.00	29.43
ATOM	173	CE1	PHE A	37	15.168	39.679	22.564	1.00	31.59
ATOM	174	CE2	PHE A	37	14.852	40.580	24.733	1.00	30.52
ATOM	175	CZ	PHE A	37	14.927	39.495	23.915	1.00	31.81
ATOM	176	N	PRO A	38	14.403	43.721	19.384	1.00	31.70
ATOM	177	CA	PRO A	38	14.268	43.036	18.106	1.00	31.23
ATOM	178	C	PRO A	38	15.231	41.895	17.984	1.00	31.71
ATOM	179	O	PRO A	38	16.205	41.880	18.743	1.00	31.59
ATOM	180	CB	PRO A	38	14.652	44.118	17.117	1.00	31.44
ATOM	181	CG	PRO A	38	15.636	45.005	17.922	1.00	32.43
ATOM	182	CD	PRO A	38	15.186	44.973	19.320	1.00	30.26
ATOM	183	N	THR A	39	14.996	41.008	17.003	1.00	31.90
ATOM	184	CA	THR A	39	15.868	39.891	16.702	1.00	32.00
ATOM	185	C	THR A	39	15.865	39.606	15.203	1.00	32.89
ATOM	186	O	THR A	39	14.998	40.048	14.476	1.00	32.28
ATOM	187	CB	THR A	39	15.370	38.647	17.377	1.00	32.26
ATOM	188	OG1	THR A	39	14.024	38.389	16.949	1.00	30.59
ATOM	189	CG2	THR A	39	15.235	38.832	18.905	1.00	32.99
ATOM	190	N	ARG A	40	16.855	38.844	14.770	1.00	33.28
ATOM	191	CA	ARG A	40	16.980	38.372	13.421	1.00	34.30
ATOM	192	C	ARG A	40	17.023	36.853	13.562	1.00	33.77
ATOM	193	O	ARG A	40	17.288	36.324	14.614	1.00	34.68
ATOM	194	CB	ARG A	40	18.267	38.847	12.795	1.00	34.73
ATOM	195	CG	ARG A	40	18.273	40.306	12.451	1.00	41.24
ATOM	196	CD	ARG A	40	16.909	40.799	12.022	1.00	48.38
ATOM	197	NE	ARG A	40	16.728	41.025	10.597	1.00	51.83
ATOM	198	CZ	ARG A	40	15.553	41.291	10.092	1.00	56.70
ATOM	199	NH1	ARG A	40	14.501	41.300	10.908	1.00	57.84
ATOM	200	NH2	ARG A	40	15.415	41.549	8.795	1.00	60.30
ATOM	201	N	PRO A	41	16.663	36.147	12.533	1.00	33.50
ATOM	202	CA	PRO A	41	16.646	34.681	12.569	1.00	33.48
ATOM	203	C	PRO A	41	18.002	34.010	12.475	1.00	32.72

C
O
O
N
C
C
O
C
C
C
C
C
O
N
C
C
C
O
N
C
C
C
C
C
C
N
C
C
C
O
C
C
C
C
N
C
C
C
C
C
C
N
C
C
N
N
C
C

ATOM	204	O	PRO	A	41	18.907	34.475	11.801	1.00	34.64
ATOM	205	CB	PRO	A	41	15.800	34.299	11.340	1.00	33.22
ATOM	206	CG	PRO	A	41	15.409	35.563	10.675	1.00	33.55
ATOM	207	CD	PRO	A	41	16.038	36.717	11.343	1.00	32.95
ATOM	208	N	ILE	A	42	18.154	32.925	13.192	1.00	30.97
ATOM	209	CA	ILE	A	42	19.337	32.144	13.049	1.00	29.87
ATOM	210	C	ILE	A	42	19.077	31.289	11.814	1.00	30.07
ATOM	211	O	ILE	A	42	17.946	30.857	11.587	1.00	30.17
ATOM	212	CB	ILE	A	42	19.477	31.306	14.273	1.00	29.85
ATOM	213	CG1	ILE	A	42	19.721	32.215	15.478	1.00	27.79
ATOM	214	CG2	ILE	A	42	20.582	30.264	14.105	1.00	30.60
ATOM	215	CD1	ILE	A	42	19.565	31.503	16.788	1.00	25.35
ATOM	216	N	PRO	A	43	20.085	31.075	10.985	1.00	29.34
ATOM	217	CA	PRO	A	43	19.922	30.219	9.823	1.00	29.60
ATOM	218	C	PRO	A	43	19.623	28.751	10.176	1.00	30.65
ATOM	219	O	PRO	A	43	20.207	28.265	11.160	1.00	31.73
ATOM	220	CB	PRO	A	43	21.288	30.310	9.139	1.00	30.00
ATOM	221	CG	PRO	A	43	21.933	31.491	9.665	1.00	28.87
ATOM	222	CD	PRO	A	43	21.420	31.676	11.047	1.00	28.76
ATOM	223	N	ARG	A	44	18.725	28.103	9.419	1.00	30.34
ATOM	224	CA	ARG	A	44	18.415	26.693	9.507	1.00	31.59
ATOM	225	C	ARG	A	44	18.965	26.050	8.267	1.00	31.44
ATOM	226	O	ARG	A	44	18.540	26.359	7.169	1.00	32.24
ATOM	227	CB	ARG	A	44	16.923	26.399	9.502	1.00	32.07
ATOM	228	CG	ARG	A	44	16.105	27.115	10.563	1.00	36.61
ATOM	229	CD	ARG	A	44	14.638	26.534	10.809	1.00	39.09
ATOM	230	NE	ARG	A	44	14.350	25.140	10.395	1.00	41.98
ATOM	231	CZ	ARG	A	44	14.179	24.099	11.246	1.00	43.25
ATOM	232	NH1	ARG	A	44	14.332	24.244	12.561	1.00	40.76
ATOM	233	NH2	ARG	A	44	13.864	22.892	10.777	1.00	44.70
ATOM	234	N	LEU	A	45	19.878	25.118	8.428	1.00	31.29
ATOM	235	CA	LEU	A	45	20.544	24.560	7.294	1.00	30.82
ATOM	236	C	LEU	A	45	20.744	23.083	7.470	1.00	31.52
ATOM	237	O	LEU	A	45	20.690	22.550	8.597	1.00	31.99
ATOM	238	CB	LEU	A	45	21.909	25.221	7.134	1.00	29.63
ATOM	239	CG	LEU	A	45	21.873	26.697	6.813	1.00	31.00
ATOM	240	CD1	LEU	A	45	23.279	27.338	7.030	1.00	31.79
ATOM	241	CD2	LEU	A	45	21.403	26.918	5.378	1.00	31.14
ATOM	242	N	SER	A	46	21.004	22.437	6.338	1.00	31.72
ATOM	243	CA	SER	A	46	21.345	21.057	6.327	1.00	32.17
ATOM	244	C	SER	A	46	22.771	20.978	6.767	1.00	32.55
ATOM	245	O	SER	A	46	23.568	21.871	6.562	1.00	31.31
ATOM	246	CB	SER	A	46	21.255	20.447	4.936	1.00	32.06
ATOM	247	OG	SER	A	46	21.909	19.187	4.931	1.00	31.15
ATOM	248	N	GLN	A	47	23.068	19.849	7.358	1.00	33.60
ATOM	249	CA	GLN	A	47	24.370	19.546	7.878	1.00	34.42
ATOM	250	C	GLN	A	47	25.385	19.453	6.754	1.00	34.74
ATOM	251	O	GLN	A	47	26.554	19.685	6.971	1.00	35.40
ATOM	252	CB	GLN	A	47	24.232	18.194	8.593	1.00	35.09
ATOM	253	CG	GLN	A	47	25.465	17.417	8.733	1.00	36.19
ATOM	254	CD	GLN	A	47	25.670	16.422	7.678	1.00	36.93
ATOM	255	OE1	GLN	A	47	24.757	16.093	6.902	1.00	38.75
ATOM	256	NE2	GLN	A	47	26.894	15.903	7.630	1.00	40.53
ATOM	257	N	SER	A	48	24.933	19.095	5.556	1.00	34.63
ATOM	258	CA	SER	A	48	25.821	18.922	4.415	1.00	34.93
ATOM	259	C	SER	A	48	26.128	20.249	3.735	1.00	35.90
ATOM	260	O	SER	A	48	26.990	20.315	2.862	1.00	36.31
ATOM	261	CB	SER	A	48	25.182	17.991	3.381	1.00	34.59
ATOM	262	OG	SER	A	48	23.912	18.483	2.955	1.00	33.48
ATOM	263	N	ASP	A	49	25.415	21.302	4.124	1.00	36.41
ATOM	264	CA	ASP	A	49	25.583	22.606	3.512	1.00	37.09

O
C
C
C
N
C
C
C
O
C
C
C
N
C
C
C
O
C
C
C
N
C
N
N
C
C
C
O
C
C
C
C
O
N
N
C
C
C
O
C
C
O
N
C

ATOM	265	C	ASP	A	49	26.855	23.291	3.989	1.00	37.81
ATOM	266	O	ASP	A	49	27.020	23.523	5.182	1.00	36.99
ATOM	267	CB	ASP	A	49	24.383	23.451	3.877	1.00	37.42
ATOM	268	CG	ASP	A	49	24.323	24.733	3.117	1.00	38.29
ATOM	269	OD1	ASP	A	49	25.383	25.351	2.849	1.00	37.80
ATOM	270	OD2	ASP	A	49	23.223	25.191	2.760	1.00	40.61
ATOM	271	N	PRO	A	50	27.740	23.649	3.056	1.00	38.39
ATOM	272	CA	PRO	A	50	29.005	24.291	3.409	1.00	38.24
ATOM	273	C	PRO	A	50	28.802	25.502	4.285	1.00	38.16
ATOM	274	O	PRO	A	50	29.671	25.797	5.088	1.00	37.37
ATOM	275	CB	PRO	A	50	29.577	24.738	2.051	1.00	38.14
ATOM	276	CG	PRO	A	50	28.978	23.915	1.070	1.00	38.31
ATOM	277	CD	PRO	A	50	27.620	23.473	1.601	1.00	38.85
ATOM	278	N	ARG	A	51	27.686	26.199	4.125	1.00	38.66
ATOM	279	CA	ARG	A	51	27.448	27.403	4.912	1.00	39.05
ATOM	280	C	ARG	A	51	27.332	27.077	6.391	1.00	38.94
ATOM	281	O	ARG	A	51	27.668	27.908	7.259	1.00	38.35
ATOM	282	CB	ARG	A	51	26.199	28.120	4.438	1.00	39.34
ATOM	283	CG	ARG	A	51	26.372	28.781	3.067	1.00	41.10
ATOM	284	CD	ARG	A	51	25.099	29.355	2.486	1.00	42.20
ATOM	285	NE	ARG	A	51	24.111	28.314	2.192	1.00	44.96
ATOM	286	CZ	ARG	A	51	22.815	28.551	2.043	1.00	46.68
ATOM	287	NH1	ARG	A	51	22.348	29.795	2.162	1.00	47.61
ATOM	288	NH2	ARG	A	51	21.980	27.563	1.774	1.00	46.08
ATOM	289	N	ALA	A	52	26.882	25.867	6.688	1.00	38.31
ATOM	290	CA	ALA	A	52	26.729	25.498	8.082	1.00	38.63
ATOM	291	C	ALA	A	52	28.093	25.311	8.677	1.00	38.81
ATOM	292	O	ALA	A	52	28.341	25.682	9.816	1.00	39.03
ATOM	293	CB	ALA	A	52	25.907	24.233	8.235	1.00	38.36
ATOM	294	N	GLU	A	53	28.988	24.733	7.899	1.00	39.48
ATOM	295	CA	GLU	A	53	30.316	24.484	8.388	1.00	40.51
ATOM	296	C	GLU	A	53	31.038	25.802	8.621	1.00	40.41
ATOM	297	O	GLU	A	53	31.740	25.954	9.618	1.00	39.82
ATOM	298	CB	GLU	A	53	31.093	23.614	7.436	1.00	40.70
ATOM	299	CG	GLU	A	53	32.129	22.790	8.171	1.00	44.79
ATOM	300	CD	GLU	A	53	31.596	21.435	8.633	1.00	48.98
ATOM	301	OE1	GLU	A	53	30.427	21.329	9.017	1.00	51.56
ATOM	302	OE2	GLU	A	53	32.354	20.454	8.620	1.00	54.44
ATOM	303	N	GLU	A	54	30.851	26.752	7.713	1.00	40.45
ATOM	304	CA	GLU	A	54	31.465	28.060	7.853	1.00	41.40
ATOM	305	C	GLU	A	54	30.991	28.731	9.162	1.00	40.32
ATOM	306	O	GLU	A	54	31.760	29.374	9.836	1.00	39.89
ATOM	307	CB	GLU	A	54	31.119	28.974	6.679	1.00	42.12
ATOM	308	CG	GLU	A	54	31.697	28.557	5.333	1.00	47.25
ATOM	309	CD	GLU	A	54	30.953	29.191	4.129	1.00	53.54
ATOM	310	OE1	GLU	A	54	30.459	30.366	4.276	1.00	57.47
ATOM	311	OE2	GLU	A	54	30.882	28.527	3.036	1.00	53.82
ATOM	312	N	LEU	A	55	29.735	28.534	9.533	1.00	39.03
ATOM	313	CA	LEU	A	55	29.223	29.198	10.687	1.00	37.80
ATOM	314	C	LEU	A	55	29.840	28.673	11.956	1.00	36.83
ATOM	315	O	LEU	A	55	30.293	29.464	12.795	1.00	35.66
ATOM	316	CB	LEU	A	55	27.715	29.081	10.725	1.00	38.06
ATOM	317	CG	LEU	A	55	27.072	29.921	9.621	1.00	37.63
ATOM	318	CD1	LEU	A	55	25.620	29.569	9.449	1.00	39.63
ATOM	319	CD2	LEU	A	55	27.174	31.356	9.928	1.00	36.27
ATOM	320	N	ILE	A	56	29.876	27.346	12.079	1.00	35.87
ATOM	321	CA	ILE	A	56	30.436	26.684	13.257	1.00	35.35
ATOM	322	C	ILE	A	56	31.916	27.001	13.410	1.00	35.82
ATOM	323	O	ILE	A	56	32.377	27.299	14.499	1.00	36.59
ATOM	324	CB	ILE	A	56	30.281	25.198	13.139	1.00	34.84
ATOM	325	CG1	ILE	A	56	28.790	24.810	13.132	1.00	34.00

C
O
C
C
O
O
N
C
C
O
C
C
N
C
C
O
C
C
N
C
N
N
N
C
C
O
C
N
C
C
C
O
C
C
C
O
O
N
C
C
C
O
C
C
C
C
N
C
C
C
O
C
C

ATOM	326	CG2	ILE	A	56	31.022	24.535	14.280	1.00	34.89
ATOM	327	CD1	ILE	A	56	28.502	23.361	12.684	1.00	30.65
ATOM	328	N	GLU	A	57	32.636	26.964	12.301	1.00	36.28
ATOM	329	CA	GLU	A	57	34.056	27.244	12.255	1.00	37.26
ATOM	330	C	GLU	A	57	34.295	28.634	12.809	1.00	37.41
ATOM	331	O	GLU	A	57	35.278	28.879	13.516	1.00	38.73
ATOM	332	CB	GLU	A	57	34.562	27.167	10.795	1.00	37.54
ATOM	333	CG	GLU	A	57	36.035	27.462	10.572	1.00	39.41
ATOM	334	CD	GLU	A	57	36.948	26.668	11.493	1.00	44.86
ATOM	335	OE1	GLU	A	57	36.666	25.458	11.741	1.00	48.72
ATOM	336	OE2	GLU	A	57	37.961	27.260	11.965	1.00	46.32
ATOM	337	N	ASN	A	58	33.386	29.537	12.485	1.00	36.64
ATOM	338	CA	ASN	A	58	33.456	30.907	12.926	1.00	36.26
ATOM	339	C	ASN	A	58	32.768	31.226	14.219	1.00	35.28
ATOM	340	O	ASN	A	58	32.569	32.376	14.531	1.00	33.45
ATOM	341	CB	ASN	A	58	32.798	31.758	11.898	1.00	36.88
ATOM	342	CG	ASN	A	58	33.763	32.461	11.107	1.00	39.77
ATOM	343	OD1	ASN	A	58	34.140	31.983	10.018	1.00	41.34
ATOM	344	ND2	ASN	A	58	34.241	33.617	11.639	1.00	41.14
ATOM	345	N	GLU	A	59	32.366	30.205	14.945	1.00	35.05
ATOM	346	CA	GLU	A	59	31.706	30.415	16.226	1.00	35.00
ATOM	347	C	GLU	A	59	30.481	31.314	16.133	1.00	33.86
ATOM	348	O	GLU	A	59	30.293	32.238	16.904	1.00	33.16
ATOM	349	CB	GLU	A	59	32.729	30.855	17.244	1.00	34.95
ATOM	350	CG	GLU	A	59	33.708	29.700	17.423	1.00	37.79
ATOM	351	CD	GLU	A	59	34.652	29.820	18.586	1.00	39.57
ATOM	352	OE1	GLU	A	59	35.809	30.145	18.349	1.00	43.67
ATOM	353	OE2	GLU	A	59	34.254	29.545	19.723	1.00	45.29
ATOM	354	N	GLU	A	60	29.644	30.995	15.157	1.00	32.92
ATOM	355	CA	GLU	A	60	28.359	31.641	14.983	1.00	32.66
ATOM	356	C	GLU	A	60	27.284	30.576	14.993	1.00	30.59
ATOM	357	O	GLU	A	60	27.456	29.485	14.461	1.00	30.47
ATOM	358	CB	GLU	A	60	28.312	32.400	13.685	1.00	33.20
ATOM	359	CG	GLU	A	60	29.384	33.454	13.615	1.00	38.62
ATOM	360	CD	GLU	A	60	28.955	34.617	12.762	1.00	45.73
ATOM	361	OE1	GLU	A	60	29.075	34.553	11.519	1.00	48.47
ATOM	362	OE2	GLU	A	60	28.466	35.587	13.367	1.00	53.89
ATOM	363	N	PRO	A	61	26.152	30.915	15.560	1.00	28.63
ATOM	364	CA	PRO	A	61	25.073	29.954	15.735	1.00	28.05
ATOM	365	C	PRO	A	61	24.462	29.451	14.443	1.00	27.61
ATOM	366	O	PRO	A	61	24.475	30.120	13.426	1.00	27.23
ATOM	367	CB	PRO	A	61	24.021	30.754	16.492	1.00	28.67
ATOM	368	CG	PRO	A	61	24.385	32.140	16.400	1.00	27.46
ATOM	369	CD	PRO	A	61	25.811	32.253	16.038	1.00	27.71
ATOM	370	N	VAL	A	62	23.936	28.237	14.475	1.00	27.43
ATOM	371	CA	VAL	A	62	23.254	27.685	13.317	1.00	27.09
ATOM	372	C	VAL	A	62	22.372	26.569	13.802	1.00	27.40
ATOM	373	O	VAL	A	62	22.707	25.893	14.780	1.00	27.79
ATOM	374	CB	VAL	A	62	24.219	27.181	12.270	1.00	27.04
ATOM	375	CG1	VAL	A	62	25.106	26.166	12.829	1.00	27.34
ATOM	376	CG2	VAL	A	62	23.473	26.659	11.037	1.00	27.58
ATOM	377	N	VAL	A	63	21.195	26.440	13.199	1.00	27.19
ATOM	378	CA	VAL	A	63	20.339	25.338	13.526	1.00	27.37
ATOM	379	C	VAL	A	63	20.540	24.324	12.383	1.00	27.33
ATOM	380	O	VAL	A	63	20.360	24.672	11.208	1.00	27.05
ATOM	381	CB	VAL	A	63	18.857	25.737	13.611	1.00	27.47
ATOM	382	CG1	VAL	A	63	17.976	24.446	13.695	1.00	28.55
ATOM	383	CG2	VAL	A	63	18.616	26.603	14.793	1.00	25.77
ATOM	384	N	LEU	A	64	20.933	23.114	12.743	1.00	27.26
ATOM	385	CA	LEU	A	64	21.119	22.001	11.820	1.00	28.51
ATOM	386	C	LEU	A	64	19.897	21.128	11.894	1.00	28.40

CCNCCOCCOONCCOCCONNCCOCCOONCCOCCOONCCOCCOCCNCCOCCNCCNCC

ATOM	387	O	LEU	A	64	19.517	20.692	12.982	1.00	28.69	O
ATOM	388	CB	LEU	A	64	22.352	21.187	12.204	1.00	28.92	C
ATOM	389	CG	LEU	A	64	23.624	22.016	12.142	1.00	31.65	C
ATOM	390	CD1	LEU	A	64	24.800	21.273	12.534	1.00	34.84	C
ATOM	391	CD2	LEU	A	64	23.843	22.490	10.722	1.00	35.77	C
ATOM	392	N	THR	A	65	19.284	20.868	10.752	1.00	28.44	N
ATOM	393	CA	THR	A	65	18.003	20.164	10.721	1.00	29.54	C
ATOM	394	C	THR	A	65	18.054	18.659	10.534	1.00	29.40	C
ATOM	395	O	THR	A	65	17.046	17.985	10.799	1.00	29.20	O
ATOM	396	CB	THR	A	65	17.139	20.679	9.552	1.00	30.07	C
ATOM	397	OG1	THR	A	65	17.879	20.569	8.318	1.00	30.75	O
ATOM	398	CG2	THR	A	65	16.878	22.137	9.665	1.00	31.86	C
ATOM	399	N	ASP	A	66	19.183	18.150	10.059	1.00	29.30	N
ATOM	400	CA	ASP	A	66	19.323	16.722	9.719	1.00	29.78	C
ATOM	401	C	ASP	A	66	20.573	15.970	10.196	1.00	29.12	C
ATOM	402	O	ASP	A	66	21.081	15.141	9.432	1.00	29.09	O
ATOM	403	CB	ASP	A	66	19.301	16.582	8.195	1.00	29.41	C
ATOM	404	CG	ASP	A	66	20.274	17.507	7.525	1.00	32.36	C
ATOM	405	OD1	ASP	A	66	20.924	18.319	8.230	1.00	34.87	O
ATOM	406	OD2	ASP	A	66	20.447	17.534	6.287	1.00	35.70	O
ATOM	407	N	THR	A	67	21.078	16.254	11.404	1.00	28.05	N
ATOM	408	CA	THR	A	67	22.266	15.565	11.897	1.00	27.24	C
ATOM	409	C	THR	A	67	21.943	14.171	12.355	1.00	26.92	C
ATOM	410	O	THR	A	67	22.807	13.297	12.367	1.00	25.98	O
ATOM	411	CB	THR	A	67	22.856	16.258	13.134	1.00	27.78	C
ATOM	412	OG1	THR	A	67	21.874	16.338	14.186	1.00	25.71	O
ATOM	413	CG2	THR	A	67	23.268	17.664	12.820	1.00	29.22	C
ATOM	414	N	ASN	A	68	20.704	13.979	12.777	1.00	26.53	N
ATOM	415	CA	ASN	A	68	20.307	12.711	13.355	1.00	27.10	C
ATOM	416	C	ASN	A	68	21.075	12.499	14.647	1.00	26.46	C
ATOM	417	O	ASN	A	68	21.223	11.399	15.092	1.00	26.90	O
ATOM	418	CB	ASN	A	68	20.583	11.547	12.401	1.00	27.41	C
ATOM	419	CG	ASN	A	68	19.546	11.415	11.308	1.00	28.18	C
ATOM	420	OD1	ASN	A	68	18.345	11.286	11.566	1.00	29.22	O
ATOM	421	ND2	ASN	A	68	20.006	11.445	10.077	1.00	29.40	N
ATOM	422	N	LEU	A	69	21.585	13.569	15.227	1.00	26.58	N
ATOM	423	CA	LEU	A	69	22.429	13.481	16.427	1.00	26.67	C
ATOM	424	C	LEU	A	69	21.832	12.639	17.558	1.00	26.73	C
ATOM	425	O	LEU	A	69	22.550	11.822	18.166	1.00	27.74	O
ATOM	426	CB	LEU	A	69	22.739	14.870	16.968	1.00	26.04	C
ATOM	427	CG	LEU	A	69	23.677	14.904	18.157	1.00	26.23	C
ATOM	428	CD1	LEU	A	69	24.986	14.168	17.894	1.00	27.92	C
ATOM	429	CD2	LEU	A	69	23.976	16.343	18.556	1.00	24.74	C
ATOM	430	N	VAL	A	70	20.557	12.857	17.861	1.00	26.02	N
ATOM	431	CA	VAL	A	70	19.882	12.091	18.898	1.00	26.50	C
ATOM	432	C	VAL	A	70	18.673	11.364	18.345	1.00	26.71	C
ATOM	433	O	VAL	A	70	17.650	11.183	19.015	1.00	25.59	O
ATOM	434	CB	VAL	A	70	19.479	12.943	20.098	1.00	26.46	C
ATOM	435	CG1	VAL	A	70	20.717	13.505	20.764	1.00	28.17	C
ATOM	436	CG2	VAL	A	70	18.517	14.016	19.716	1.00	23.88	C
ATOM	437	N	TYR	A	71	18.832	10.920	17.113	1.00	27.86	N
ATOM	438	CA	TYR	A	71	17.791	10.159	16.451	1.00	29.60	C
ATOM	439	C	TYR	A	71	17.093	9.167	17.388	1.00	29.76	C
ATOM	440	O	TYR	A	71	15.905	9.202	17.478	1.00	30.54	O
ATOM	441	CB	TYR	A	71	18.326	9.459	15.208	1.00	29.31	C
ATOM	442	CG	TYR	A	71	17.347	8.479	14.686	1.00	31.42	C
ATOM	443	CD1	TYR	A	71	16.136	8.884	14.120	1.00	31.39	C
ATOM	444	CD2	TYR	A	71	17.610	7.107	14.779	1.00	32.30	C
ATOM	445	CE1	TYR	A	71	15.223	7.905	13.641	1.00	31.86	C
ATOM	446	CE2	TYR	A	71	16.736	6.181	14.317	1.00	29.75	C
ATOM	447	CZ	TYR	A	71	15.556	6.558	13.755	1.00	31.10	C

ATOM	448	OH	TYR	A	71	14.698	5.530	13.342	1.00	39.44
ATOM	449	N	PRO	A	72	17.807	8.317	18.108	1.00	30.56
ATOM	450	CA	PRO	A	72	17.150	7.336	18.991	1.00	30.85
ATOM	451	C	PRO	A	72	16.389	7.914	20.158	1.00	30.85
ATOM	452	O	PRO	A	72	15.549	7.233	20.750	1.00	30.12
ATOM	453	CB	PRO	A	72	18.310	6.468	19.507	1.00	30.71
ATOM	454	CG	PRO	A	72	19.420	6.692	18.515	1.00	31.79
ATOM	455	CD	PRO	A	72	19.268	8.158	18.096	1.00	30.88
ATOM	456	N	ALA	A	73	16.650	9.166	20.508	1.00	31.72
ATOM	457	CA	ALA	A	73	15.918	9.743	21.627	1.00	31.71
ATOM	458	C	ALA	A	73	14.596	10.371	21.172	1.00	31.83
ATOM	459	O	ALA	A	73	13.786	10.808	22.000	1.00	31.52
ATOM	460	CB	ALA	A	73	16.762	10.742	22.340	1.00	32.17
ATOM	461	N	LEU	A	74	14.345	10.408	19.868	1.00	31.19
ATOM	462	CA	LEU	A	74	13.108	11.028	19.439	1.00	32.01
ATOM	463	C	LEU	A	74	11.856	10.343	19.965	1.00	32.30
ATOM	464	O	LEU	A	74	10.807	10.914	19.963	1.00	31.90
ATOM	465	CB	LEU	A	74	13.050	11.191	17.931	1.00	31.25
ATOM	466	CG	LEU	A	74	14.140	12.103	17.400	1.00	34.53
ATOM	467	CD1	LEU	A	74	13.904	12.418	15.965	1.00	37.35
ATOM	468	CD2	LEU	A	74	14.264	13.436	18.200	1.00	34.44
ATOM	469	N	LYS	A	75	11.963	9.100	20.396	1.00	34.18
ATOM	470	CA	LYS	A	75	10.802	8.376	20.910	1.00	34.36
ATOM	471	C	LYS	A	75	10.618	8.616	22.403	1.00	35.18
ATOM	472	O	LYS	A	75	9.575	8.304	22.945	1.00	35.70
ATOM	473	CB	LYS	A	75	10.950	6.876	20.631	1.00	34.51
ATOM	474	CG	LYS	A	75	12.138	6.219	21.294	1.00	33.19
ATOM	475	CD	LYS	A	75	12.302	4.754	20.834	1.00	31.54
ATOM	476	CE	LYS	A	75	13.796	4.304	20.744	1.00	29.70
ATOM	477	NZ	LYS	A	75	14.460	4.158	22.026	1.00	28.79
ATOM	478	N	TRP	A	76	11.619	9.181	23.071	1.00	35.44
ATOM	479	CA	TRP	A	76	11.517	9.426	24.495	1.00	36.05
ATOM	480	C	TRP	A	76	10.307	10.264	24.893	1.00	37.47
ATOM	481	O	TRP	A	76	9.958	11.222	24.212	1.00	38.43
ATOM	482	CB	TRP	A	76	12.742	10.179	24.994	1.00	35.67
ATOM	483	CG	TRP	A	76	13.980	9.395	24.963	1.00	33.87
ATOM	484	CD1	TRP	A	76	14.136	8.116	24.520	1.00	32.29
ATOM	485	CD2	TRP	A	76	15.252	9.821	25.409	1.00	30.36
ATOM	486	NE1	TRP	A	76	15.440	7.723	24.673	1.00	30.42
ATOM	487	CE2	TRP	A	76	16.143	8.754	25.221	1.00	28.82
ATOM	488	CE3	TRP	A	76	15.739	11.013	25.954	1.00	29.45
ATOM	489	CZ2	TRP	A	76	17.471	8.833	25.560	1.00	30.56
ATOM	490	CZ3	TRP	A	76	17.056	11.091	26.291	1.00	29.56
ATOM	491	CH2	TRP	A	76	17.917	10.017	26.092	1.00	30.20
ATOM	492	N	ASP	A	77	9.698	9.894	26.014	1.00	38.14
ATOM	493	CA	ASP	A	77	8.620	10.640	26.635	1.00	38.33
ATOM	494	C	ASP	A	77	8.653	10.223	28.094	1.00	37.89
ATOM	495	O	ASP	A	77	9.443	9.390	28.447	1.00	37.97
ATOM	496	CB	ASP	A	77	7.283	10.353	25.974	1.00	38.71
ATOM	497	CG	ASP	A	77	6.882	8.904	26.049	1.00	39.50
ATOM	498	OD1	ASP	A	77	7.414	8.107	26.867	1.00	41.50
ATOM	499	OD2	ASP	A	77	6.006	8.472	25.299	1.00	41.98
ATOM	500	N	LEU	A	78	7.838	10.800	28.955	1.00	38.47
ATOM	501	CA	LEU	A	78	7.941	10.481	30.375	1.00	39.09
ATOM	502	C	LEU	A	78	7.734	9.002	30.662	1.00	39.89
ATOM	503	O	LEU	A	78	8.421	8.423	31.507	1.00	40.19
ATOM	504	CB	LEU	A	78	6.966	11.310	31.171	1.00	39.02
ATOM	505	CG	LEU	A	78	7.169	12.815	31.042	1.00	41.51
ATOM	506	CD1	LEU	A	78	6.058	13.567	31.764	1.00	42.72
ATOM	507	CD2	LEU	A	78	8.522	13.216	31.618	1.00	42.31
ATOM	508	N	GLU	A	79	6.795	8.369	29.971	1.00	40.66

ATOM	509	CA	GLU	A	79	6.557	6.948	30.212	1.00	41.43	C
ATOM	510	C	GLU	A	79	7.794	6.086	29.909	1.00	40.84	C
ATOM	511	O	GLU	A	79	8.228	5.283	30.742	1.00	40.45	O
ATOM	512	CB	GLU	A	79	5.343	6.458	29.418	1.00	42.10	C
ATOM	513	CG	GLU	A	79	5.023	4.998	29.658	1.00	45.06	C
ATOM	514	CD	GLU	A	79	3.840	4.525	28.846	1.00	49.74	C
ATOM	515	OE1	GLU	A	79	3.436	5.254	27.914	1.00	52.37	O
ATOM	516	OE2	GLU	A	79	3.316	3.426	29.150	1.00	53.21	O
ATOM	517	N	TYR	A	80	8.361	6.248	28.718	1.00	40.43	N
ATOM	518	CA	TYR	A	80	9.559	5.495	28.362	1.00	40.14	C
ATOM	519	C	TYR	A	80	10.750	5.765	29.297	1.00	40.06	C
ATOM	520	O	TYR	A	80	11.485	4.844	29.664	1.00	39.72	O
ATOM	521	CB	TYR	A	80	9.946	5.832	26.946	1.00	39.96	C
ATOM	522	CG	TYR	A	80	11.193	5.147	26.420	1.00	39.51	C
ATOM	523	CD1	TYR	A	80	11.118	3.897	25.804	1.00	38.93	C
ATOM	524	CD2	TYR	A	80	12.433	5.768	26.490	1.00	36.89	C
ATOM	525	CE1	TYR	A	80	12.253	3.277	25.290	1.00	38.08	C
ATOM	526	CE2	TYR	A	80	13.562	5.154	25.993	1.00	38.15	C
ATOM	527	CZ	TYR	A	80	13.468	3.915	25.379	1.00	39.06	C
ATOM	528	OH	TYR	A	80	14.604	3.319	24.863	1.00	38.99	O
ATOM	529	N	LEU	A	81	10.935	7.017	29.688	1.00	39.68	N
ATOM	530	CA	LEU	A	81	12.061	7.362	30.565	1.00	40.01	C
ATOM	531	C	LEU	A	81	11.858	6.806	31.978	1.00	39.86	C
ATOM	532	O	LEU	A	81	12.792	6.284	32.608	1.00	39.47	O
ATOM	533	CB	LEU	A	81	12.314	8.889	30.604	1.00	39.50	C
ATOM	534	CG	LEU	A	81	12.765	9.532	29.287	1.00	39.76	C
ATOM	535	CD1	LEU	A	81	12.805	11.045	29.402	1.00	39.84	C
ATOM	536	CD2	LEU	A	81	14.117	9.018	28.840	1.00	39.32	C
ATOM	537	N	GLN	A	82	10.645	6.932	32.482	1.00	40.38	N
ATOM	538	CA	GLN	A	82	10.346	6.415	33.804	1.00	41.02	C
ATOM	539	C	GLN	A	82	10.612	4.920	33.799	1.00	40.84	C
ATOM	540	O	GLN	A	82	11.193	4.379	34.711	1.00	40.35	O
ATOM	541	CB	GLN	A	82	8.900	6.688	34.164	1.00	41.40	C
ATOM	542	CG	GLN	A	82	8.447	5.906	35.351	1.00	42.97	C
ATOM	543	CD	GLN	A	82	7.291	6.560	36.099	1.00	45.14	C
ATOM	544	OE1	GLN	A	82	6.662	7.502	35.616	1.00	44.10	O
ATOM	545	NE2	GLN	A	82	7.003	6.039	37.287	1.00	48.04	N
ATOM	546	N	GLU	A	83	10.222	4.256	32.728	1.00	41.35	N
ATOM	547	CA	GLU	A	83	10.442	2.816	32.620	1.00	41.46	C
ATOM	548	C	GLU	A	83	11.926	2.444	32.465	1.00	40.81	C
ATOM	549	O	GLU	A	83	12.350	1.407	32.958	1.00	41.13	O
ATOM	550	CB	GLU	A	83	9.610	2.243	31.472	1.00	41.27	C
ATOM	551	CG	GLU	A	83	9.609	0.728	31.366	1.00	44.77	C
ATOM	552	CD	GLU	A	83	8.878	0.018	32.514	1.00	49.16	C
ATOM	553	OE1	GLU	A	83	8.219	0.672	33.347	1.00	52.26	O
ATOM	554	OE2	GLU	A	83	8.978	-1.220	32.606	1.00	53.23	O
ATOM	555	N	ASN	A	84	12.743	3.298	31.854	1.00	39.81	N
ATOM	556	CA	ASN	A	84	14.099	2.859	31.515	1.00	38.82	C
ATOM	557	C	ASN	A	84	15.290	3.611	32.048	1.00	38.20	C
ATOM	558	O	ASN	A	84	16.402	3.117	31.919	1.00	36.98	O
ATOM	559	CB	ASN	A	84	14.244	2.828	29.995	1.00	38.83	C
ATOM	560	CG	ASN	A	84	13.301	1.865	29.349	1.00	38.45	C
ATOM	561	OD1	ASN	A	84	13.307	0.670	29.664	1.00	40.40	O
ATOM	562	ND2	ASN	A	84	12.480	2.365	28.441	1.00	35.77	N
ATOM	563	N	ILE	A	85	15.088	4.781	32.645	1.00	38.08	N
ATOM	564	CA	ILE	A	85	16.226	5.598	32.998	1.00	38.57	C
ATOM	565	C	ILE	A	85	16.929	5.280	34.307	1.00	38.70	C
ATOM	566	O	ILE	A	85	17.849	5.993	34.702	1.00	39.78	O
ATOM	567	CB	ILE	A	85	15.827	7.056	32.932	1.00	39.02	C
ATOM	568	CG1	ILE	A	85	17.001	7.891	32.418	1.00	39.43	C
ATOM	569	CG2	ILE	A	85	15.291	7.544	34.266	1.00	39.22	C

ATOM	631	O	SER A	94	11.851	26.181	42.441	1.00	43.63
ATOM	632	CB	SER A	94	10.570	23.415	43.051	1.00	42.76
ATOM	633	OG	SER A	94	9.258	23.841	43.395	1.00	45.05
ATOM	634	N	ALA A	95	9.929	26.171	41.308	1.00	43.42
ATOM	635	CA	ALA A	95	9.839	27.607	41.323	1.00	43.49
ATOM	636	C	ALA A	95	8.416	28.083	41.281	1.00	43.87
ATOM	637	O	ALA A	95	7.520	27.416	40.789	1.00	43.95
ATOM	638	CB	ALA A	95	10.572	28.164	40.145	1.00	43.70
ATOM	639	N	SER A	96	8.219	29.287	41.770	1.00	44.66
ATOM	640	CA	SER A	96	6.901	29.866	41.757	1.00	45.08
ATOM	641	C	SER A	96	6.767	30.914	40.668	1.00	44.27
ATOM	642	O	SER A	96	5.783	31.636	40.637	1.00	45.20
ATOM	643	CB	SER A	96	6.604	30.510	43.095	1.00	45.33
ATOM	644	OG	SER A	96	5.384	31.199	42.973	1.00	47.96
ATOM	645	N	THR A	97	7.776	31.034	39.819	1.00	42.95
ATOM	646	CA	THR A	97	7.738	31.962	38.688	1.00	42.08
ATOM	647	C	THR A	97	8.103	31.184	37.461	1.00	40.94
ATOM	648	O	THR A	97	8.604	30.095	37.584	1.00	40.73
ATOM	649	CB	THR A	97	8.779	33.088	38.826	1.00	42.48
ATOM	650	OG1	THR A	97	8.964	33.707	37.546	1.00	43.60
ATOM	651	CG2	THR A	97	10.209	32.544	39.156	1.00	42.12
ATOM	652	N	HIS A	98	7.904	31.734	36.275	1.00	40.18
ATOM	653	CA	HIS A	98	8.319	31.012	35.074	1.00	40.09
ATOM	654	C	HIS A	98	9.840	31.036	34.883	1.00	39.78
ATOM	655	O	HIS A	98	10.376	30.277	34.078	1.00	38.48
ATOM	656	CB	HIS A	98	7.660	31.580	33.824	1.00	40.00
ATOM	657	CG	HIS A	98	7.947	33.028	33.582	1.00	40.56
ATOM	658	ND1	HIS A	98	7.297	34.041	34.259	1.00	40.13
ATOM	659	CD2	HIS A	98	8.805	33.640	32.729	1.00	42.26
ATOM	660	CE1	HIS A	98	7.741	35.212	33.841	1.00	37.08
ATOM	661	NE2	HIS A	98	8.661	35.002	32.915	1.00	39.74
ATOM	662	N	LYS A	99	10.526	31.883	35.652	1.00	39.73
ATOM	663	CA	LYS A	99	11.948	32.082	35.476	1.00	40.45
ATOM	664	C	LYS A	99	12.861	31.221	36.332	1.00	40.34
ATOM	665	O	LYS A	99	12.904	31.376	37.554	1.00	41.93
ATOM	666	CB	LYS A	99	12.279	33.535	35.730	1.00	40.89
ATOM	667	CG	LYS A	99	11.819	34.443	34.624	1.00	43.12
ATOM	668	CD	LYS A	99	12.466	35.806	34.753	1.00	46.04
ATOM	669	CE	LYS A	99	11.733	36.727	35.711	1.00	48.26
ATOM	670	NZ	LYS A	99	10.662	37.495	35.003	1.00	49.55
ATOM	671	N	PHE A	100	13.622	30.335	35.700	1.00	39.70
ATOM	672	CA	PHE A	100	14.554	29.487	36.447	1.00	39.48
ATOM	673	C	PHE A	100	16.001	30.011	36.390	1.00	39.52
ATOM	674	O	PHE A	100	16.874	29.435	35.718	1.00	38.87
ATOM	675	CB	PHE A	100	14.485	28.056	35.939	1.00	39.33
ATOM	676	CG	PHE A	100	13.266	27.328	36.377	1.00	38.47
ATOM	677	CD1	PHE A	100	12.059	27.519	35.725	1.00	36.20
ATOM	678	CD2	PHE A	100	13.327	26.441	37.443	1.00	37.31
ATOM	679	CE1	PHE A	100	10.960	26.846	36.125	1.00	36.52
ATOM	680	CE2	PHE A	100	12.212	25.751	37.849	1.00	36.40
ATOM	681	CZ	PHE A	100	11.037	25.946	37.206	1.00	36.52
ATOM	682	N	LEU A	101	16.244	31.130	37.066	1.00	39.53
ATOM	683	CA	LEU A	101	17.592	31.667	37.170	1.00	39.36
ATOM	684	C	LEU A	101	18.548	30.642	37.787	1.00	39.88
ATOM	685	O	LEU A	101	18.345	30.152	38.899	1.00	39.47
ATOM	686	CB	LEU A	101	17.574	32.912	38.036	1.00	39.37
ATOM	687	CG	LEU A	101	18.828	33.777	38.071	1.00	39.44
ATOM	688	CD1	LEU A	101	19.258	34.290	36.693	1.00	37.56
ATOM	689	CD2	LEU A	101	18.546	34.926	39.008	1.00	39.90
ATOM	690	N	TYR A	102	19.578	30.297	37.032	1.00	41.10
ATOM	691	CA	TYR A	102	20.623	29.415	37.507	1.00	41.59

ATOM	692	C	TYR A 102	21.439	30.097	38.611	1.00	42.23
ATOM	693	O	TYR A 102	21.778	31.296	38.500	1.00	42.28
ATOM	694	CB	TYR A 102	21.580	29.048	36.370	1.00	41.49
ATOM	695	CG	TYR A 102	22.706	28.209	36.894	1.00	43.02
ATOM	696	CD1	TYR A 102	22.507	26.864	37.200	1.00	42.85
ATOM	697	CD2	TYR A 102	23.941	28.761	37.150	1.00	43.85
ATOM	698	CE1	TYR A 102	23.504	26.110	37.725	1.00	45.04
ATOM	699	CE2	TYR A 102	24.942	28.005	37.696	1.00	46.05
ATOM	700	CZ	TYR A 102	24.725	26.681	37.984	1.00	46.73
ATOM	701	OH	TYR A 102	25.752	25.915	38.544	1.00	52.49
ATOM	702	N	TYR A 103	21.759	29.340	39.657	1.00	42.12
ATOM	703	CA	TYR A 103	22.665	29.832	40.690	1.00	42.58
ATOM	704	C	TYR A 103	23.632	28.750	41.207	1.00	41.98
ATOM	705	O	TYR A 103	23.333	27.560	41.237	1.00	42.05
ATOM	706	CB	TYR A 103	21.901	30.435	41.843	1.00	43.03
ATOM	707	CG	TYR A 103	20.964	29.493	42.477	1.00	44.79
ATOM	708	CD1	TYR A 103	19.691	29.333	41.971	1.00	47.69
ATOM	709	CD2	TYR A 103	21.338	28.769	43.590	1.00	49.51
ATOM	710	CE1	TYR A 103	18.798	28.456	42.544	1.00	50.63
ATOM	711	CE2	TYR A 103	20.449	27.890	44.202	1.00	52.43
ATOM	712	CZ	TYR A 103	19.181	27.736	43.658	1.00	52.68
ATOM	713	OH	TYR A 103	18.297	26.875	44.234	1.00	56.40
ATOM	714	N	ASP A 104	24.809	29.190	41.603	1.00	41.00
ATOM	715	CA	ASP A 104	25.844	28.288	42.026	1.00	40.33
ATOM	716	C	ASP A 104	25.842	28.250	43.530	1.00	39.98
ATOM	717	O	ASP A 104	26.177	29.224	44.219	1.00	38.58
ATOM	718	CB	ASP A 104	27.169	28.768	41.492	1.00	40.13
ATOM	719	CG	ASP A 104	28.310	27.929	41.958	1.00	42.26
ATOM	720	OD1	ASP A 104	28.117	27.151	42.942	1.00	44.33
ATOM	721	OD2	ASP A 104	29.429	27.989	41.393	1.00	42.30
ATOM	722	N	GLU A 105	25.425	27.110	44.041	1.00	40.20
ATOM	723	CA	GLU A 105	25.294	26.950	45.472	1.00	40.26
ATOM	724	C	GLU A 105	26.605	27.213	46.228	1.00	39.92
ATOM	725	O	GLU A 105	26.577	27.824	47.293	1.00	40.11
ATOM	726	CB	GLU A 105	24.747	25.570	45.744	1.00	40.54
ATOM	727	CG	GLU A 105	23.245	25.502	45.533	1.00	42.55
ATOM	728	CD	GLU A 105	22.709	24.086	45.506	1.00	44.11
ATOM	729	OE1	GLU A 105	23.178	23.251	46.323	1.00	46.78
ATOM	730	OE2	GLU A 105	21.801	23.815	44.684	1.00	42.74
ATOM	731	N	LYS A 106	27.748	26.819	45.661	1.00	39.52
ATOM	732	CA	LYS A 106	29.041	26.991	46.336	1.00	39.53
ATOM	733	C	LYS A 106	29.370	28.419	46.623	1.00	39.68
ATOM	734	O	LYS A 106	30.199	28.697	47.471	1.00	40.24
ATOM	735	CB	LYS A 106	30.221	26.479	45.504	1.00	39.89
ATOM	736	CG	LYS A 106	30.257	24.976	45.143	1.00	40.25
ATOM	737	N	LYS A 107	28.747	29.333	45.894	1.00	40.07
ATOM	738	CA	LYS A 107	29.043	30.740	46.031	1.00	39.35
ATOM	739	C	LYS A 107	28.061	31.421	46.937	1.00	39.82
ATOM	740	O	LYS A 107	28.208	32.586	47.188	1.00	39.28
ATOM	741	CB	LYS A 107	29.050	31.410	44.645	1.00	39.29
ATOM	742	CG	LYS A 107	30.329	31.155	43.827	1.00	37.47
ATOM	743	CD	LYS A 107	30.286	31.702	42.406	1.00	34.55
ATOM	744	CE	LYS A 107	31.606	31.470	41.651	1.00	33.57
ATOM	745	NZ	LYS A 107	31.607	31.880	40.178	1.00	30.95
ATOM	746	N	MET A 108	27.071	30.707	47.458	1.00	41.78
ATOM	747	CA	MET A 108	26.079	31.341	48.341	1.00	43.69
ATOM	748	C	MET A 108	26.621	31.950	49.646	1.00	44.68
ATOM	749	O	MET A 108	26.154	33.004	50.061	1.00	44.99
ATOM	750	CB	MET A 108	24.923	30.396	48.594	1.00	44.05
ATOM	751	CG	MET A 108	24.195	30.105	47.282	1.00	46.49
ATOM	752	SD	MET A 108	22.723	29.125	47.315	1.00	51.54

[illegible]

ATOM	753	CE	MET	A	108	21.653	30.080	48.484	1.00	51.43
ATOM	754	N	ALA	A	109	27.630	31.341	50.254	1.00	46.34
ATOM	755	CA	ALA	A	109	28.211	31.855	51.510	1.00	47.87
ATOM	756	C	ALA	A	109	28.697	33.308	51.406	1.00	49.27
ATOM	757	O	ALA	A	109	28.533	34.114	52.323	1.00	49.44
ATOM	758	CB	ALA	A	109	29.355	30.958	51.960	1.00	47.63
ATOM	759	N	ASN	A	110	29.276	33.651	50.269	1.00	50.88
ATOM	760	CA	ASN	A	110	29.792	35.004	50.069	1.00	51.99
ATOM	761	C	ASN	A	110	28.732	36.070	49.736	1.00	51.88
ATOM	762	O	ASN	A	110	29.038	37.252	49.562	1.00	51.14
ATOM	763	CB	ASN	A	110	30.876	34.939	48.994	1.00	52.43
ATOM	764	CG	ASN	A	110	32.077	34.080	49.433	1.00	54.41
ATOM	765	OD1	ASN	A	110	32.460	34.050	50.624	1.00	54.72
ATOM	766	ND2	ASN	A	110	32.663	33.373	48.478	1.00	56.75
ATOM	767	N	PHE	A	111	27.484	35.651	49.632	1.00	52.59
ATOM	768	CA	PHE	A	111	26.404	36.598	49.409	1.00	53.32
ATOM	769	C	PHE	A	111	25.202	36.180	50.261	1.00	54.75
ATOM	770	O	PHE	A	111	24.149	35.801	49.749	1.00	54.37
ATOM	771	CB	PHE	A	111	26.053	36.693	47.923	1.00	52.83
ATOM	772	CG	PHE	A	111	27.058	37.461	47.107	1.00	50.42
ATOM	773	CD1	PHE	A	111	28.283	36.907	46.791	1.00	49.93
ATOM	774	CD2	PHE	A	111	26.781	38.729	46.658	1.00	48.60
ATOM	775	CE1	PHE	A	111	29.205	37.609	46.044	1.00	49.18
ATOM	776	CE2	PHE	A	111	27.701	39.429	45.904	1.00	47.91
ATOM	777	CZ	PHE	A	111	28.910	38.869	45.601	1.00	47.42
ATOM	778	N	GLN	A	112	25.376	36.279	51.576	1.00	56.74
ATOM	779	CA	GLN	A	112	24.353	35.855	52.536	1.00	58.14
ATOM	780	C	GLN	A	112	22.970	36.399	52.192	1.00	58.29
ATOM	781	O	GLN	A	112	21.972	35.742	52.468	1.00	58.53
ATOM	782	CB	GLN	A	112	24.730	36.282	53.961	1.00	58.65
ATOM	783	CG	GLN	A	112	26.100	35.815	54.441	1.00	61.33
ATOM	784	CD	GLN	A	112	26.213	34.307	54.467	1.00	64.34
ATOM	785	OE1	GLN	A	112	25.483	33.622	53.750	1.00	66.46
ATOM	786	NE2	GLN	A	112	27.133	33.784	55.281	1.00	66.00
ATOM	787	N	ASN	A	113	22.919	37.581	51.580	1.00	58.45
ATOM	788	CA	ASN	A	113	21.653	38.222	51.240	1.00	58.79
ATOM	789	C	ASN	A	113	21.000	37.740	49.936	1.00	58.82
ATOM	790	O	ASN	A	113	19.971	38.284	49.536	1.00	59.69
ATOM	791	CB	ASN	A	113	21.813	39.759	51.171	1.00	58.81
ATOM	792	CG	ASN	A	113	22.160	40.387	52.525	1.00	59.17
ATOM	793	OD1	ASN	A	113	21.863	39.824	53.593	1.00	57.49
ATOM	794	ND2	ASN	A	113	22.786	41.567	52.482	1.00	58.34
ATOM	795	N	PHE	A	114	21.570	36.760	49.245	1.00	58.33
ATOM	796	CA	PHE	A	114	20.930	36.306	48.015	1.00	57.84
ATOM	797	C	PHE	A	114	19.922	35.220	48.334	1.00	57.57
ATOM	798	O	PHE	A	114	20.264	34.208	48.953	1.00	57.24
ATOM	799	CB	PHE	A	114	21.933	35.774	47.013	1.00	57.99
ATOM	800	CG	PHE	A	114	21.289	35.126	45.822	1.00	58.28
ATOM	801	CD1	PHE	A	114	20.642	35.897	44.867	1.00	58.11
ATOM	802	CD2	PHE	A	114	21.300	33.744	45.674	1.00	58.56
ATOM	803	CE1	PHE	A	114	20.028	35.312	43.781	1.00	58.29
ATOM	804	CE2	PHE	A	114	20.687	33.149	44.580	1.00	59.09
ATOM	805	CZ	PHE	A	114	20.050	33.937	43.630	1.00	58.80
ATOM	806	N	LYS	A	115	18.686	35.429	47.899	1.00	57.18
ATOM	807	CA	LYS	A	115	17.604	34.511	48.204	1.00	57.19
ATOM	808	C	LYS	A	115	17.037	33.942	46.903	1.00	57.17
ATOM	809	O	LYS	A	115	16.268	34.598	46.193	1.00	57.12
ATOM	810	CB	LYS	A	115	16.524	35.234	49.016	1.00	57.33
ATOM	811	N	PRO	A	116	17.384	32.699	46.610	1.00	56.97
ATOM	812	CA	PRO	A	116	17.042	32.097	45.319	1.00	57.01
ATOM	813	C	PRO	A	116	15.554	31.979	45.151	1.00	56.69

C
N
C
C
O
C
N
C
C
O
C
O
N
N
C
C
O
C
C
C
C
C
C
N
C
C
C
O
C
C
O
C
C
C
C
C
N
C
C
C
O
C
C
N
C
C

ATOM	814	O	PRO	A	116	14.864	31.676	46.103	1.00	57.03	O
ATOM	815	CB	PRO	A	116	17.672	30.701	45.378	1.00	57.15	C
ATOM	816	CG	PRO	A	116	18.474	30.658	46.662	1.00	57.53	C
ATOM	817	CD	PRO	A	116	18.045	31.763	47.522	1.00	57.03	C
ATOM	818	N	ARG	A	117	15.073	32.212	43.945	1.00	56.65	N
ATOM	819	CA	ARG	A	117	13.663	32.087	43.656	1.00	56.39	C
ATOM	820	C	ARG	A	117	13.335	30.639	43.332	1.00	56.16	C
ATOM	821	O	ARG	A	117	12.160	30.272	43.279	1.00	56.65	O
ATOM	822	CB	ARG	A	117	13.280	32.971	42.508	1.00	56.56	C
ATOM	823	N	SER	A	118	14.350	29.810	43.110	1.00	55.25	N
ATOM	824	CA	SER	A	118	14.083	28.393	42.878	1.00	55.12	C
ATOM	825	C	SER	A	118	14.924	27.511	43.809	1.00	54.45	C
ATOM	826	O	SER	A	118	16.022	27.895	44.174	1.00	54.64	O
ATOM	827	CB	SER	A	118	14.322	28.029	41.415	1.00	54.89	C
ATOM	828	OG	SER	A	118	15.672	28.196	41.087	1.00	55.13	O
ATOM	829	N	ASN	A	119	14.381	26.362	44.211	1.00	53.66	N
ATOM	830	CA	ASN	A	119	15.089	25.412	45.060	1.00	53.43	C
ATOM	831	C	ASN	A	119	15.307	24.118	44.326	1.00	51.73	C
ATOM	832	O	ASN	A	119	14.472	23.705	43.552	1.00	51.23	O
ATOM	833	CB	ASN	A	119	14.277	25.063	46.311	1.00	54.32	C
ATOM	834	CG	ASN	A	119	13.711	26.272	46.994	1.00	57.18	C
ATOM	835	OD1	ASN	A	119	14.448	27.099	47.563	1.00	60.84	O
ATOM	836	ND2	ASN	A	119	12.384	26.384	46.964	1.00	61.01	N
ATOM	837	N	ARG	A	120	16.418	23.470	44.621	1.00	50.77	N
ATOM	838	CA	ARG	A	120	16.772	22.182	44.049	1.00	49.89	C
ATOM	839	C	ARG	A	120	16.388	21.062	44.999	1.00	49.87	C
ATOM	840	O	ARG	A	120	16.620	21.150	46.206	1.00	50.40	O
ATOM	841	CB	ARG	A	120	18.283	22.133	43.843	1.00	49.75	C
ATOM	842	CG	ARG	A	120	18.799	20.901	43.144	1.00	47.86	C
ATOM	843	CD	ARG	A	120	20.318	20.812	43.074	1.00	45.29	C
ATOM	844	NE	ARG	A	120	20.972	21.240	44.301	1.00	43.79	N
ATOM	845	CZ	ARG	A	120	21.316	20.432	45.314	1.00	43.05	C
ATOM	846	NH1	ARG	A	120	21.053	19.126	45.277	1.00	40.07	N
ATOM	847	NH2	ARG	A	120	21.921	20.942	46.379	1.00	41.03	N
ATOM	848	N	GLU	A	121	15.801	20.010	44.451	1.00	49.51	N
ATOM	849	CA	GLU	A	121	15.435	18.830	45.206	1.00	48.90	C
ATOM	850	C	GLU	A	121	15.905	17.609	44.411	1.00	47.95	C
ATOM	851	O	GLU	A	121	15.793	17.564	43.192	1.00	48.02	O
ATOM	852	CB	GLU	A	121	13.922	18.782	45.461	1.00	49.38	C
ATOM	853	CG	GLU	A	121	13.444	17.500	46.141	1.00	51.77	C
ATOM	854	CD	GLU	A	121	12.185	17.695	46.961	1.00	54.65	C
ATOM	855	OE1	GLU	A	121	12.312	18.258	48.073	1.00	58.68	O
ATOM	856	OE2	GLU	A	121	11.087	17.284	46.512	1.00	55.59	O
ATOM	857	N	GLU	A	122	16.455	16.622	45.104	1.00	47.05	N
ATOM	858	CA	GLU	A	122	16.938	15.423	44.463	1.00	46.05	C
ATOM	859	C	GLU	A	122	15.848	14.400	44.594	1.00	45.48	C
ATOM	860	O	GLU	A	122	15.313	14.220	45.666	1.00	45.63	O
ATOM	861	CB	GLU	A	122	18.200	14.940	45.158	1.00	45.67	C
ATOM	862	CG	GLU	A	122	19.346	15.921	45.063	1.00	44.97	C
ATOM	863	CD	GLU	A	122	19.789	16.154	43.623	1.00	45.49	C
ATOM	864	OE1	GLU	A	122	19.985	15.144	42.902	1.00	44.79	O
ATOM	865	OE2	GLU	A	122	19.925	17.337	43.207	1.00	42.47	O
ATOM	866	N	MET	A	123	15.481	13.751	43.503	1.00	44.95	N
ATOM	867	CA	MET	A	123	14.462	12.731	43.589	1.00	44.35	C
ATOM	868	C	MET	A	123	14.657	11.712	42.503	1.00	43.59	C
ATOM	869	O	MET	A	123	15.495	11.891	41.613	1.00	43.44	O
ATOM	870	CB	MET	A	123	13.078	13.358	43.490	1.00	44.59	C
ATOM	871	CG	MET	A	123	12.831	14.092	42.205	1.00	45.95	C
ATOM	872	SD	MET	A	123	11.244	14.946	42.206	1.00	47.99	S
ATOM	873	CE	MET	A	123	11.648	16.411	43.089	1.00	47.81	C
ATOM	874	N	LYS	A	124	13.880	10.638	42.603	1.00	42.37	N

ATOM	875	CA	LYS	A	124	13.869	9.575	41.623	1.00	41.47	C
ATOM	876	C	LYS	A	124	12.958	10.026	40.469	1.00	40.66	C
ATOM	877	O	LYS	A	124	12.066	10.862	40.654	1.00	39.80	O
ATOM	878	CB	LYS	A	124	13.354	8.267	42.250	1.00	41.45	C
ATOM	879	CG	LYS	A	124	14.160	7.756	43.470	1.00	41.70	C
ATOM	880	CD	LYS	A	124	15.574	7.301	43.076	1.00	42.20	C
ATOM	881	CE	LYS	A	124	16.451	6.989	44.273	1.00	41.37	C
ATOM	882	NZ	LYS	A	124	17.894	7.201	43.966	1.00	41.63	N
ATOM	883	N	PHE	A	125	13.178	9.460	39.284	1.00	39.69	N
ATOM	884	CA	PHE	A	125	12.463	9.889	38.101	1.00	38.95	C
ATOM	885	C	PHE	A	125	10.981	9.729	38.257	1.00	38.85	C
ATOM	886	O	PHE	A	125	10.217	10.613	37.890	1.00	37.81	O
ATOM	887	CB	PHE	A	125	12.942	9.154	36.862	1.00	38.45	C
ATOM	888	CG	PHE	A	125	12.618	9.871	35.610	1.00	39.49	C
ATOM	889	CD1	PHE	A	125	13.471	10.840	35.118	1.00	40.78	C
ATOM	890	CD2	PHE	A	125	11.428	9.655	34.967	1.00	40.81	C
ATOM	891	CE1	PHE	A	125	13.181	11.530	33.992	1.00	40.48	C
ATOM	892	CE2	PHE	A	125	11.123	10.353	33.825	1.00	41.39	C
ATOM	893	CZ	PHE	A	125	12.008	11.300	33.335	1.00	41.11	C
ATOM	894	N	HIS	A	126	10.575	8.602	38.835	1.00	39.49	N
ATOM	895	CA	HIS	A	126	9.161	8.342	39.044	1.00	39.91	C
ATOM	896	C	HIS	A	126	8.599	9.408	39.976	1.00	39.98	C
ATOM	897	O	HIS	A	126	7.455	9.821	39.830	1.00	39.26	O
ATOM	898	CB	HIS	A	126	8.906	6.899	39.552	1.00	40.45	C
ATOM	899	CG	HIS	A	126	9.034	6.727	41.033	1.00	41.06	C
ATOM	900	ND1	HIS	A	126	7.987	6.950	41.899	1.00	40.98	N
ATOM	901	CD2	HIS	A	126	10.092	6.367	41.801	1.00	41.92	C
ATOM	902	CE1	HIS	A	126	8.403	6.759	43.140	1.00	43.22	C
ATOM	903	NE2	HIS	A	126	9.676	6.405	43.108	1.00	42.45	N
ATOM	904	N	GLU	A	127	9.428	9.898	40.890	1.00	40.59	N
ATOM	905	CA	GLU	A	127	8.975	10.919	41.824	1.00	41.56	C
ATOM	906	C	GLU	A	127	8.749	12.213	41.062	1.00	41.66	C
ATOM	907	O	GLU	A	127	7.774	12.941	41.294	1.00	41.64	O
ATOM	908	CB	GLU	A	127	9.968	11.102	42.978	1.00	41.64	C
ATOM	909	CG	GLU	A	127	10.149	9.839	43.811	1.00	43.10	C
ATOM	910	CD	GLU	A	127	11.116	10.017	44.958	1.00	44.79	C
ATOM	911	OE1	GLU	A	127	12.326	10.217	44.712	1.00	43.71	O
ATOM	912	OE2	GLU	A	127	10.649	9.938	46.122	1.00	49.43	O
ATOM	913	N	PHE	A	128	9.636	12.491	40.124	1.00	41.96	N
ATOM	914	CA	PHE	A	128	9.504	13.705	39.337	1.00	41.99	C
ATOM	915	C	PHE	A	128	8.213	13.612	38.538	1.00	42.56	C
ATOM	916	O	PHE	A	128	7.400	14.537	38.507	1.00	41.93	O
ATOM	917	CB	PHE	A	128	10.708	13.864	38.416	1.00	41.57	C
ATOM	918	CG	PHE	A	128	10.470	14.795	37.244	1.00	41.38	C
ATOM	919	CD1	PHE	A	128	10.199	16.136	37.449	1.00	39.65	C
ATOM	920	CD2	PHE	A	128	10.537	14.330	35.950	1.00	39.22	C
ATOM	921	CE1	PHE	A	128	9.980	16.975	36.399	1.00	40.36	C
ATOM	922	CE2	PHE	A	128	10.329	15.191	34.888	1.00	41.57	C
ATOM	923	CZ	PHE	A	128	10.044	16.510	35.112	1.00	40.14	C
ATOM	924	N	VAL	A	129	7.999	12.463	37.921	1.00	43.85	N
ATOM	925	CA	VAL	A	129	6.813	12.288	37.095	1.00	44.87	C
ATOM	926	C	VAL	A	129	5.536	12.443	37.932	1.00	46.07	C
ATOM	927	O	VAL	A	129	4.574	13.093	37.505	1.00	45.32	O
ATOM	928	CB	VAL	A	129	6.798	10.920	36.425	1.00	44.99	C
ATOM	929	CG1	VAL	A	129	5.496	10.725	35.631	1.00	44.81	C
ATOM	930	CG2	VAL	A	129	8.030	10.718	35.550	1.00	43.96	C
ATOM	931	N	GLU	A	130	5.553	11.854	39.128	1.00	47.64	N
ATOM	932	CA	GLU	A	130	4.415	11.932	40.037	1.00	49.12	C
ATOM	933	C	GLU	A	130	4.136	13.390	40.384	1.00	49.48	C
ATOM	934	O	GLU	A	130	3.003	13.846	40.250	1.00	48.66	O
ATOM	935	CB	GLU	A	130	4.654	11.079	41.296	1.00	49.63	C

ATOM	936	CG	GLU	A	130	4.534	9.576	41.062	1.00	50.81
ATOM	937	CD	GLU	A	130	5.257	8.747	42.117	1.00	53.77
ATOM	938	OE1	GLU	A	130	5.566	9.295	43.204	1.00	55.27
ATOM	939	OE2	GLU	A	130	5.518	7.543	41.856	1.00	54.73
ATOM	940	N	LYS	A	131	5.170	14.121	40.798	1.00	50.56
ATOM	941	CA	LYS	A	131	5.010	15.547	41.082	1.00	51.84
ATOM	942	C	LYS	A	131	4.385	16.276	39.886	1.00	52.55
ATOM	943	O	LYS	A	131	3.482	17.070	40.061	1.00	52.15
ATOM	944	CB	LYS	A	131	6.340	16.209	41.438	1.00	52.23
ATOM	945	CG	LYS	A	131	6.554	16.568	42.907	1.00	53.75
ATOM	946	CD	LYS	A	131	7.249	17.944	43.054	1.00	56.27
ATOM	947	CE	LYS	A	131	7.879	18.168	44.448	1.00	58.66
ATOM	948	NZ	LYS	A	131	8.247	19.628	44.725	1.00	59.31
ATOM	949	N	LEU	A	132	4.845	16.014	38.667	1.00	53.84
ATOM	950	CA	LEU	A	132	4.247	16.691	37.513	1.00	55.23
ATOM	951	C	LEU	A	132	2.762	16.408	37.407	1.00	55.96
ATOM	952	O	LEU	A	132	1.973	17.292	37.063	1.00	55.69
ATOM	953	CB	LEU	A	132	4.906	16.278	36.206	1.00	55.45
ATOM	954	CG	LEU	A	132	6.301	16.811	35.966	1.00	56.60
ATOM	955	CD1	LEU	A	132	6.785	16.291	34.646	1.00	57.60
ATOM	956	CD2	LEU	A	132	6.303	18.327	35.966	1.00	57.89
ATOM	957	N	GLN	A	133	2.397	15.157	37.669	1.00	57.19
ATOM	958	CA	GLN	A	133	0.998	14.739	37.663	1.00	58.15
ATOM	959	C	GLN	A	133	0.172	15.450	38.751	1.00	58.99
ATOM	960	O	GLN	A	133	-0.915	15.950	38.470	1.00	58.73
ATOM	961	CB	GLN	A	133	0.897	13.230	37.824	1.00	58.03
ATOM	962	N	ASP	A	134	0.678	15.510	39.981	1.00	60.11
ATOM	963	CA	ASP	A	134	-0.081	16.145	41.063	1.00	61.49
ATOM	964	C	ASP	A	134	-0.362	17.629	40.783	1.00	61.55
ATOM	965	O	ASP	A	134	-1.427	18.149	41.104	1.00	61.55
ATOM	966	CB	ASP	A	134	0.621	15.985	42.417	1.00	61.99
ATOM	967	CG	ASP	A	134	-0.253	16.454	43.587	1.00	64.74
ATOM	968	OD1	ASP	A	134	-1.310	17.082	43.346	1.00	67.52
ATOM	969	OD2	ASP	A	134	0.022	16.237	44.791	1.00	69.05
ATOM	970	N	ILE	A	135	0.594	18.312	40.179	1.00	61.80
ATOM	971	CA	ILE	A	135	0.402	19.706	39.860	1.00	62.08
ATOM	972	C	ILE	A	135	-0.700	19.856	38.823	1.00	62.29
ATOM	973	O	ILE	A	135	-1.639	20.620	39.013	1.00	61.91
ATOM	974	CB	ILE	A	135	1.711	20.301	39.366	1.00	62.21
ATOM	975	CG1	ILE	A	135	2.705	20.391	40.528	1.00	62.47
ATOM	976	CG2	ILE	A	135	1.478	21.670	38.783	1.00	62.19
ATOM	977	CD1	ILE	A	135	4.157	20.437	40.100	1.00	62.96
ATOM	978	N	GLN	A	136	-0.593	19.112	37.733	1.00	62.99
ATOM	979	CA	GLN	A	136	-1.585	19.184	36.674	1.00	63.71
ATOM	980	C	GLN	A	136	-2.977	18.992	37.259	1.00	64.44
ATOM	981	O	GLN	A	136	-3.844	19.856	37.123	1.00	64.73
ATOM	982	CB	GLN	A	136	-1.305	18.136	35.619	1.00	63.77
ATOM	983	N	GLN	A	137	-3.170	17.871	37.948	1.00	65.12
ATOM	984	CA	GLN	A	137	-4.472	17.509	38.497	1.00	65.60
ATOM	985	C	GLN	A	137	-4.989	18.540	39.477	1.00	65.70
ATOM	986	O	GLN	A	137	-6.175	18.871	39.447	1.00	66.17
ATOM	987	CB	GLN	A	137	-4.421	16.117	39.161	1.00	65.72
ATOM	988	N	ARG	A	138	-4.115	19.056	40.336	1.00	65.61
ATOM	989	CA	ARG	A	138	-4.549	20.011	41.358	1.00	65.53
ATOM	990	C	ARG	A	138	-4.578	21.441	40.823	1.00	64.96
ATOM	991	O	ARG	A	138	-4.748	22.392	41.589	1.00	64.93
ATOM	992	CB	ARG	A	138	-3.670	19.912	42.613	1.00	65.65
ATOM	993	CG	ARG	A	138	-2.375	20.712	42.588	1.00	67.10
ATOM	994	CD	ARG	A	138	-1.482	20.352	43.745	1.00	68.96
ATOM	995	NE	ARG	A	138	-0.372	21.270	43.957	1.00	69.79
ATOM	996	CZ	ARG	A	138	0.905	20.919	43.880	1.00	71.69

C
C
O
O
N
C
C
O
C
C
C
N
N
C
C
O
C
C
N
C
C
O
C
C
O
O
N
C
C
O
C
C
C
N
C
C
O
C
C
N
C
C
O
C
C
C
N
C

[illegible]

ATOM	1058	C	LEU	A	146	15.753	18.977	34.864	1.00	35.24
ATOM	1059	O	LEU	A	146	15.665	19.211	33.653	1.00	34.63
ATOM	1060	CB	LEU	A	146	14.008	17.312	35.358	1.00	35.79
ATOM	1061	CG	LEU	A	146	14.998	16.157	35.614	1.00	37.21
ATOM	1062	CD1	LEU	A	146	14.221	14.918	35.969	1.00	36.28
ATOM	1063	CD2	LEU	A	146	15.907	15.872	34.399	1.00	38.78
ATOM	1064	N	GLN	A	147	16.907	18.883	35.515	1.00	34.82
ATOM	1065	CA	GLN	A	147	18.179	19.206	34.884	1.00	35.00
ATOM	1066	C	GLN	A	147	19.222	18.341	35.525	1.00	34.90
ATOM	1067	O	GLN	A	147	19.626	18.568	36.664	1.00	35.65
ATOM	1068	CB	GLN	A	147	18.526	20.679	35.078	1.00	35.09
ATOM	1069	CG	GLN	A	147	17.392	21.636	34.720	1.00	35.83
ATOM	1070	CD	GLN	A	147	17.848	23.085	34.706	1.00	36.90
ATOM	1071	OE1	GLN	A	147	17.091	23.961	34.302	1.00	39.61
ATOM	1072	NE2	GLN	A	147	19.088	23.336	35.126	1.00	35.05
ATOM	1073	N	GLN	A	148	19.673	17.358	34.773	1.00	35.00
ATOM	1074	CA	GLN	A	148	20.492	16.300	35.294	1.00	34.89
ATOM	1075	C	GLN	A	148	21.376	15.709	34.255	1.00	35.35
ATOM	1076	O	GLN	A	148	20.941	15.318	33.181	1.00	35.25
ATOM	1077	CB	GLN	A	148	19.580	15.186	35.791	1.00	35.27
ATOM	1078	CG	GLN	A	148	20.335	13.914	36.269	1.00	35.11
ATOM	1079	CD	GLN	A	148	21.317	14.230	37.389	1.00	34.12
ATOM	1080	OE1	GLN	A	148	20.998	15.033	38.277	1.00	33.18
ATOM	1081	NE2	GLN	A	148	22.521	13.652	37.324	1.00	30.47
ATOM	1082	N	THR	A	149	22.631	15.641	34.600	1.00	36.56
ATOM	1083	CA	THR	A	149	23.648	15.088	33.754	1.00	38.26
ATOM	1084	C	THR	A	149	23.394	13.609	33.615	1.00	38.31
ATOM	1085	O	THR	A	149	23.068	12.956	34.597	1.00	38.70
ATOM	1086	CB	THR	A	149	24.992	15.362	34.463	1.00	38.76
ATOM	1087	OG1	THR	A	149	25.290	16.747	34.277	1.00	41.68
ATOM	1088	CG2	THR	A	149	26.139	14.710	33.797	1.00	40.77
ATOM	1089	N	LEU	A	150	23.516	13.087	32.399	1.00	38.77
ATOM	1090	CA	LEU	A	150	23.373	11.664	32.149	1.00	39.27
ATOM	1091	C	LEU	A	150	24.604	10.898	32.679	1.00	39.85
ATOM	1092	O	LEU	A	150	25.733	11.207	32.321	1.00	39.99
ATOM	1093	CB	LEU	A	150	23.220	11.402	30.660	1.00	39.15
ATOM	1094	CG	LEU	A	150	21.943	11.860	29.962	1.00	40.72
ATOM	1095	CD1	LEU	A	150	22.058	11.673	28.459	1.00	41.91
ATOM	1096	CD2	LEU	A	150	20.764	11.106	30.458	1.00	41.30
ATOM	1097	N	ASN	A	151	24.392	9.898	33.526	1.00	40.26
ATOM	1098	CA	ASN	A	151	25.510	9.124	34.062	1.00	40.90
ATOM	1099	C	ASN	A	151	25.291	7.607	34.041	1.00	41.95
ATOM	1100	O	ASN	A	151	24.327	7.098	33.441	1.00	41.72
ATOM	1101	CB	ASN	A	151	25.778	9.575	35.489	1.00	40.77
ATOM	1102	CG	ASN	A	151	24.564	9.439	36.349	1.00	39.85
ATOM	1103	OD1	ASN	A	151	24.002	8.340	36.483	1.00	37.92
ATOM	1104	ND2	ASN	A	151	24.107	10.558	36.897	1.00	36.89
ATOM	1105	N	ASP	A	152	26.176	6.894	34.735	1.00	42.89
ATOM	1106	CA	ASP	A	152	26.175	5.431	34.756	1.00	43.75
ATOM	1107	C	ASP	A	152	24.950	4.768	35.272	1.00	43.37
ATOM	1108	O	ASP	A	152	24.787	3.581	35.052	1.00	43.66
ATOM	1109	CB	ASP	A	152	27.256	4.895	35.695	1.00	44.77
ATOM	1110	CG	ASP	A	152	28.539	5.555	35.500	1.00	48.22
ATOM	1111	OD1	ASP	A	152	28.835	5.881	34.325	1.00	57.47
ATOM	1112	OD2	ASP	A	152	29.291	5.826	36.437	1.00	51.28
ATOM	1113	N	THR	A	153	24.119	5.466	36.025	1.00	43.41
ATOM	1114	CA	THR	A	153	22.983	4.783	36.631	1.00	43.38
ATOM	1115	C	THR	A	153	21.820	4.605	35.676	1.00	42.99
ATOM	1116	O	THR	A	153	20.866	3.945	36.012	1.00	43.44
ATOM	1117	CB	THR	A	153	22.489	5.507	37.875	1.00	43.44
ATOM	1118	OG1	THR	A	153	21.749	6.678	37.499	1.00	46.09

C
O
C
C
C
C
N
C
C
O
C
C
O
N
N
C
C
C
O
C
C
O
N
N
C
C
C
O
C
C
C
N
C
C
C
O
C
C
O
N
N
C
C
C
O
C
C
O
O
O
N
C
C
C
O
C
O

ATOM	1119	CG2	THR	A	153	23.651	6.008	38.725	1.00	42.99
ATOM	1120	N	VAL	A	154	21.874	5.185	34.491	1.00	42.88
ATOM	1121	CA	VAL	A	154	20.757	5.015	33.572	1.00	42.71
ATOM	1122	C	VAL	A	154	20.700	3.563	33.177	1.00	42.42
ATOM	1123	O	VAL	A	154	21.729	2.892	33.193	1.00	42.39
ATOM	1124	CB	VAL	A	154	20.889	5.875	32.300	1.00	42.67
ATOM	1125	CG1	VAL	A	154	20.857	7.347	32.660	1.00	43.04
ATOM	1126	CG2	VAL	A	154	22.159	5.516	31.538	1.00	42.56
ATOM	1127	N	GLY	A	155	19.502	3.097	32.814	1.00	42.11
ATOM	1128	CA	GLY	A	155	19.261	1.725	32.411	1.00	41.52
ATOM	1129	C	GLY	A	155	19.642	1.317	30.992	1.00	41.85
ATOM	1130	O	GLY	A	155	19.977	2.140	30.123	1.00	41.87
ATOM	1131	N	ARG	A	156	19.512	0.012	30.750	1.00	41.36
ATOM	1132	CA	ARG	A	156	19.961	-0.623	29.511	1.00	40.88
ATOM	1133	C	ARG	A	156	19.397	-0.005	28.247	1.00	40.03
ATOM	1134	O	ARG	A	156	20.135	0.304	27.339	1.00	40.36
ATOM	1135	CB	ARG	A	156	19.670	-2.148	29.555	1.00	40.77
ATOM	1136	N	LYS	A	157	18.090	0.146	28.181	1.00	39.16
ATOM	1137	CA	LYS	A	157	17.503	0.697	26.990	1.00	38.81
ATOM	1138	C	LYS	A	157	18.078	2.114	26.738	1.00	38.38
ATOM	1139	O	LYS	A	157	18.363	2.464	25.594	1.00	37.97
ATOM	1140	CB	LYS	A	157	15.968	0.714	27.090	1.00	39.20
ATOM	1141	CG	LYS	A	157	15.266	-0.575	26.642	1.00	36.80
ATOM	1142	N	ILE	A	158	18.268	2.909	27.791	1.00	37.56
ATOM	1143	CA	ILE	A	158	18.759	4.287	27.606	1.00	37.28
ATOM	1144	C	ILE	A	158	20.180	4.281	27.126	1.00	37.22
ATOM	1145	O	ILE	A	158	20.582	5.120	26.302	1.00	37.08
ATOM	1146	CB	ILE	A	158	18.692	5.101	28.882	1.00	36.72
ATOM	1147	CG1	ILE	A	158	17.257	5.220	29.356	1.00	37.17
ATOM	1148	CG2	ILE	A	158	19.254	6.465	28.646	1.00	36.48
ATOM	1149	CD1	ILE	A	158	16.359	5.900	28.409	1.00	38.28
ATOM	1150	N	VAL	A	159	20.934	3.335	27.655	1.00	36.85
ATOM	1151	CA	VAL	A	159	22.319	3.177	27.284	1.00	37.05
ATOM	1152	C	VAL	A	159	22.349	2.864	25.785	1.00	36.45
ATOM	1153	O	VAL	A	159	23.165	3.410	25.057	1.00	35.81
ATOM	1154	CB	VAL	A	159	22.988	2.042	28.096	1.00	37.36
ATOM	1155	CG1	VAL	A	159	24.248	1.567	27.427	1.00	38.43
ATOM	1156	CG2	VAL	A	159	23.302	2.499	29.494	1.00	37.81
ATOM	1157	N	MET	A	160	21.453	1.990	25.339	1.00	35.84
ATOM	1158	CA	MET	A	160	21.369	1.637	23.910	1.00	36.30
ATOM	1159	C	MET	A	160	21.064	2.883	23.074	1.00	34.46
ATOM	1160	O	MET	A	160	21.715	3.142	22.083	1.00	33.88
ATOM	1161	CB	MET	A	160	20.294	0.576	23.656	1.00	36.83
ATOM	1162	CG	MET	A	160	20.654	-0.763	24.225	1.00	39.89
ATOM	1163	SD	MET	A	160	21.916	-1.579	23.249	1.00	46.87
ATOM	1164	CE	MET	A	160	21.005	-1.940	21.669	1.00	47.08
ATOM	1165	N	ASP	A	161	20.078	3.648	23.505	1.00	33.13
ATOM	1166	CA	ASP	A	161	19.732	4.901	22.857	1.00	32.80
ATOM	1167	C	ASP	A	161	20.931	5.877	22.747	1.00	31.75
ATOM	1168	O	ASP	A	161	21.209	6.425	21.691	1.00	31.06
ATOM	1169	CB	ASP	A	161	18.598	5.553	23.631	1.00	32.93
ATOM	1170	CG	ASP	A	161	17.278	4.822	23.473	1.00	33.90
ATOM	1171	OD1	ASP	A	161	17.148	3.961	22.568	1.00	28.71
ATOM	1172	OD2	ASP	A	161	16.301	5.064	24.226	1.00	38.17
ATOM	1173	N	PHE	A	162	21.646	6.052	23.849	1.00	31.05
ATOM	1174	CA	PHE	A	162	22.763	6.968	23.930	1.00	30.52
ATOM	1175	C	PHE	A	162	23.835	6.553	22.979	1.00	28.89
ATOM	1176	O	PHE	A	162	24.477	7.369	22.371	1.00	29.43
ATOM	1177	CB	PHE	A	162	23.311	6.948	25.375	1.00	30.80
ATOM	1178	CG	PHE	A	162	24.493	7.839	25.617	1.00	31.81
ATOM	1179	CD1	PHE	A	162	24.325	9.179	25.904	1.00	35.81

C
N
C
C
C
O
C
C
C
N
C
C
O
N
C
C
O
C
N
C
C
C
O
C
C
C
N
C
C
C
O
C
C
S
C
N
C
C
O
C
C
O
O
N
C
C
C
O
C
C

ATOM	1180	CD2	PHE	A	162	25.770	7.322	25.653	1.00	35.00
ATOM	1181	CE1	PHE	A	162	25.430	10.000	26.197	1.00	35.94
ATOM	1182	CE2	PHE	A	162	26.887	8.143	25.941	1.00	35.90
ATOM	1183	CZ	PHE	A	162	26.711	9.467	26.202	1.00	35.61
ATOM	1184	N	LEU	A	163	24.067	5.271	22.914	1.00	28.19
ATOM	1185	CA	LEU	A	163	25.103	4.728	22.065	1.00	28.08
ATOM	1186	C	LEU	A	163	24.735	4.923	20.590	1.00	27.13
ATOM	1187	O	LEU	A	163	25.603	4.973	19.751	1.00	26.83
ATOM	1188	CB	LEU	A	163	25.306	3.244	22.368	1.00	27.86
ATOM	1189	CG	LEU	A	163	26.137	2.987	23.604	1.00	28.73
ATOM	1190	CD1	LEU	A	163	26.180	1.512	23.849	1.00	29.77
ATOM	1191	CD2	LEU	A	163	27.559	3.522	23.464	1.00	29.70
ATOM	1192	N	GLY	A	164	23.448	4.998	20.303	1.00	26.50
ATOM	1193	CA	GLY	A	164	22.956	5.244	18.959	1.00	27.37
ATOM	1194	C	GLY	A	164	22.949	6.712	18.493	1.00	27.88
ATOM	1195	O	GLY	A	164	22.483	6.988	17.365	1.00	28.42
ATOM	1196	N	PHE	A	165	23.420	7.635	19.344	1.00	26.36
ATOM	1197	CA	PHE	A	165	23.530	9.011	18.958	1.00	26.57
ATOM	1198	C	PHE	A	165	24.540	9.052	17.820	1.00	26.12
ATOM	1199	O	PHE	A	165	25.381	8.202	17.756	1.00	24.23
ATOM	1200	CB	PHE	A	165	24.024	9.867	20.153	1.00	26.86
ATOM	1201	CG	PHE	A	165	22.979	10.070	21.252	1.00	27.80
ATOM	1202	CD1	PHE	A	165	21.701	9.533	21.144	1.00	27.80
ATOM	1203	CD2	PHE	A	165	23.274	10.820	22.367	1.00	28.58
ATOM	1204	CE1	PHE	A	165	20.765	9.744	22.117	1.00	26.89
ATOM	1205	CE2	PHE	A	165	22.329	11.042	23.364	1.00	28.45
ATOM	1206	CZ	PHE	A	165	21.086	10.504	23.243	1.00	28.78
ATOM	1207	N	ASN	A	166	24.496	10.077	16.963	1.00	26.15
ATOM	1208	CA	ASN	A	166	25.441	10.165	15.837	1.00	25.91
ATOM	1209	C	ASN	A	166	26.861	10.675	16.226	1.00	26.12
ATOM	1210	O	ASN	A	166	27.288	11.793	15.894	1.00	25.48
ATOM	1211	CB	ASN	A	166	24.834	11.004	14.705	1.00	25.51
ATOM	1212	CG	ASN	A	166	25.634	10.916	13.431	1.00	24.43
ATOM	1213	OD1	ASN	A	166	26.751	10.323	13.429	1.00	23.40
ATOM	1214	ND2	ASN	A	166	25.078	11.479	12.318	1.00	17.22
ATOM	1215	N	TRP	A	167	27.566	9.817	16.940	1.00	26.11
ATOM	1216	CA	TRP	A	167	28.907	10.080	17.401	1.00	26.58
ATOM	1217	C	TRP	A	167	29.817	10.336	16.206	1.00	27.07
ATOM	1218	O	TRP	A	167	30.737	11.139	16.252	1.00	26.92
ATOM	1219	CB	TRP	A	167	29.389	8.860	18.249	1.00	26.02
ATOM	1220	CG	TRP	A	167	28.537	8.750	19.498	1.00	27.18
ATOM	1221	CD1	TRP	A	167	27.620	7.784	19.809	1.00	28.08
ATOM	1222	CD2	TRP	A	167	28.452	9.719	20.550	1.00	27.33
ATOM	1223	NE1	TRP	A	167	27.013	8.071	21.013	1.00	27.87
ATOM	1224	CE2	TRP	A	167	27.503	9.254	21.481	1.00	26.46
ATOM	1225	CE3	TRP	A	167	29.103	10.929	20.804	1.00	26.22
ATOM	1226	CZ2	TRP	A	167	27.180	9.951	22.622	1.00	28.27
ATOM	1227	CZ3	TRP	A	167	28.784	11.615	21.920	1.00	27.56
ATOM	1228	CH2	TRP	A	167	27.821	11.128	22.833	1.00	28.04
ATOM	1229	N	ASN	A	168	29.585	9.632	15.114	1.00	27.98
ATOM	1230	CA	ASN	A	168	30.430	9.833	13.959	1.00	28.32
ATOM	1231	C	ASN	A	168	30.470	11.319	13.573	1.00	29.00
ATOM	1232	O	ASN	A	168	31.547	11.894	13.390	1.00	30.41
ATOM	1233	CB	ASN	A	168	29.909	9.018	12.812	1.00	28.58
ATOM	1234	CG	ASN	A	168	30.758	9.145	11.570	1.00	29.28
ATOM	1235	OD1	ASN	A	168	31.899	8.739	11.579	1.00	30.27
ATOM	1236	ND2	ASN	A	168	30.186	9.677	10.487	1.00	28.46
ATOM	1237	N	TRP	A	169	29.301	11.942	13.462	1.00	27.93
ATOM	1238	CA	TRP	A	169	29.228	13.343	13.074	1.00	27.25
ATOM	1239	C	TRP	A	169	29.726	14.266	14.179	1.00	27.20
ATOM	1240	O	TRP	A	169	30.444	15.224	13.925	1.00	26.30

C
C
C
C
N
C
C
O
C
C
C
N
C
C
O
N
C
C
O
C
C
C
C
N
C
C
O
C
C
C
O
N
N
C
C
O
C
C
C
C
N
C
C
C
O
C
C
O
N
N
C
C
O

ATOM	1241	CB	TRP	A	169	27.777	13.734	12.691	1.00	26.63	C
ATOM	1242	CG	TRP	A	169	27.615	15.168	12.355	1.00	25.15	C
ATOM	1243	CD1	TRP	A	169	27.804	15.745	11.130	1.00	24.89	C
ATOM	1244	CD2	TRP	A	169	27.261	16.229	13.237	1.00	23.66	C
ATOM	1245	NE1	TRP	A	169	27.591	17.095	11.195	1.00	22.98	N
ATOM	1246	CE2	TRP	A	169	27.257	17.427	12.476	1.00	25.81	C
ATOM	1247	CE3	TRP	A	169	26.927	16.292	14.576	1.00	24.44	C
ATOM	1248	CZ2	TRP	A	169	26.933	18.680	13.018	1.00	26.96	C
ATOM	1249	CZ3	TRP	A	169	26.624	17.525	15.127	1.00	28.68	C
ATOM	1250	CH2	TRP	A	169	26.611	18.710	14.336	1.00	28.73	C
ATOM	1251	N	ILE	A	170	29.323	14.026	15.409	1.00	27.29	N
ATOM	1252	CA	ILE	A	170	29.725	14.985	16.406	1.00	28.36	C
ATOM	1253	C	ILE	A	170	31.252	14.874	16.686	1.00	29.15	C
ATOM	1254	O	ILE	A	170	31.920	15.875	16.866	1.00	29.81	O
ATOM	1255	CB	ILE	A	170	28.814	14.947	17.672	1.00	28.10	C
ATOM	1256	CG1	ILE	A	170	28.882	16.277	18.386	1.00	26.98	C
ATOM	1257	CG2	ILE	A	170	29.198	13.838	18.593	1.00	27.15	C
ATOM	1258	CD1	ILE	A	170	27.855	16.400	19.530	1.00	28.88	C
ATOM	1259	N	ASN	A	171	31.809	13.679	16.653	1.00	29.35	N
ATOM	1260	CA	ASN	A	171	33.261	13.535	16.843	1.00	30.18	C
ATOM	1261	C	ASN	A	171	34.060	14.259	15.752	1.00	30.67	C
ATOM	1262	O	ASN	A	171	35.117	14.823	16.036	1.00	31.32	O
ATOM	1263	CB	ASN	A	171	33.705	12.058	16.861	1.00	29.80	C
ATOM	1264	CG	ASN	A	171	33.210	11.274	18.094	1.00	30.17	C
ATOM	1265	OD1	ASN	A	171	32.682	11.821	19.051	1.00	30.65	O
ATOM	1266	ND2	ASN	A	171	33.372	9.968	18.036	1.00	31.44	N
ATOM	1267	N	LYS	A	172	33.591	14.231	14.509	1.00	30.94	N
ATOM	1268	CA	LYS	A	172	34.297	14.960	13.461	1.00	32.29	C
ATOM	1269	C	LYS	A	172	34.211	16.473	13.755	1.00	31.79	C
ATOM	1270	O	LYS	A	172	35.128	17.234	13.482	1.00	32.06	O
ATOM	1271	CB	LYS	A	172	33.754	14.646	12.055	1.00	32.47	C
ATOM	1272	CG	LYS	A	172	34.168	13.298	11.539	1.00	36.36	C
ATOM	1273	CD	LYS	A	172	33.697	13.012	10.072	1.00	41.05	C
ATOM	1274	CE	LYS	A	172	34.338	11.721	9.543	1.00	44.14	C
ATOM	1275	NZ	LYS	A	172	34.321	11.571	8.024	1.00	49.11	N
ATOM	1276	N	GLN	A	173	33.099	16.900	14.316	1.00	31.59	N
ATOM	1277	CA	GLN	A	173	32.931	18.299	14.616	1.00	31.80	C
ATOM	1278	C	GLN	A	173	33.965	18.680	15.681	1.00	32.04	C
ATOM	1279	O	GLN	A	173	34.670	19.687	15.534	1.00	32.67	O
ATOM	1280	CB	GLN	A	173	31.522	18.581	15.117	1.00	31.45	C
ATOM	1281	CG	GLN	A	173	30.477	18.685	14.067	1.00	31.68	C
ATOM	1282	CD	GLN	A	173	30.782	19.751	13.050	1.00	32.76	C
ATOM	1283	OE1	GLN	A	173	31.198	20.852	13.405	1.00	36.55	O
ATOM	1284	NE2	GLN	A	173	30.583	19.440	11.790	1.00	32.28	N
ATOM	1285	N	GLN	A	174	34.044	17.871	16.735	1.00	31.40	N
ATOM	1286	CA	GLN	A	174	34.999	18.074	17.811	1.00	31.28	C
ATOM	1287	C	GLN	A	174	36.437	18.098	17.244	1.00	30.94	C
ATOM	1288	O	GLN	A	174	37.253	18.939	17.602	1.00	30.50	O
ATOM	1289	CB	GLN	A	174	34.808	16.971	18.861	1.00	31.20	C
ATOM	1290	CG	GLN	A	174	35.859	16.917	19.935	1.00	31.64	C
ATOM	1291	CD	GLN	A	174	35.704	15.717	20.836	1.00	31.52	C
ATOM	1292	OE1	GLN	A	174	35.313	14.661	20.382	1.00	35.10	O
ATOM	1293	NE2	GLN	A	174	36.007	15.880	22.121	1.00	32.92	N
ATOM	1294	N	GLY	A	175	36.732	17.202	16.317	1.00	30.54	N
ATOM	1295	CA	GLY	A	175	38.048	17.169	15.721	1.00	30.38	C
ATOM	1296	C	GLY	A	175	38.288	18.339	14.783	1.00	31.21	C
ATOM	1297	O	GLY	A	175	39.339	18.974	14.812	1.00	31.31	O
ATOM	1298	N	LYS	A	176	37.317	18.661	13.946	1.00	31.67	N
ATOM	1299	CA	LYS	A	176	37.524	19.767	13.029	1.00	32.84	C
ATOM	1300	C	LYS	A	176	37.758	21.127	13.749	1.00	32.90	C
ATOM	1301	O	LYS	A	176	38.563	21.906	13.291	1.00	32.85	O

ATOM	1302	CB	LYS	A	176	36.351	19.915	12.074	1.00	33.80	C
ATOM	1303	CG	LYS	A	176	36.187	18.832	11.037	1.00	35.93	C
ATOM	1304	CD	LYS	A	176	34.982	19.185	10.168	1.00	39.44	C
ATOM	1305	CE	LYS	A	176	34.055	18.032	9.961	1.00	40.94	C
ATOM	1306	NZ	LYS	A	176	32.686	18.495	9.578	1.00	45.01	N
ATOM	1307	N	ARG	A	177	37.074	21.398	14.859	1.00	32.98	N
ATOM	1308	CA	ARG	A	177	37.242	22.677	15.571	1.00	33.61	C
ATOM	1309	C	ARG	A	177	38.313	22.713	16.678	1.00	33.16	C
ATOM	1310	O	ARG	A	177	38.453	23.717	17.370	1.00	33.01	O
ATOM	1311	CB	ARG	A	177	35.920	23.101	16.220	1.00	33.86	C
ATOM	1312	CG	ARG	A	177	34.716	22.947	15.353	1.00	35.87	C
ATOM	1313	CD	ARG	A	177	34.728	23.721	14.062	1.00	37.56	C
ATOM	1314	NE	ARG	A	177	33.704	23.135	13.222	1.00	42.96	N
ATOM	1315	CZ	ARG	A	177	33.803	22.893	11.928	1.00	45.93	C
ATOM	1316	NH1	ARG	A	177	34.897	23.218	11.247	1.00	46.65	N
ATOM	1317	NH2	ARG	A	177	32.770	22.349	11.304	1.00	47.47	N
ATOM	1318	N	GLY	A	178	39.038	21.624	16.887	1.00	32.79	N
ATOM	1319	CA	GLY	A	178	40.068	21.629	17.907	1.00	31.78	C
ATOM	1320	C	GLY	A	178	39.511	21.614	19.313	1.00	31.64	C
ATOM	1321	O	GLY	A	178	40.251	21.782	20.292	1.00	33.25	O
ATOM	1322	N	TRP	A	179	38.223	21.380	19.468	1.00	30.56	N
ATOM	1323	CA	TRP	A	179	37.690	21.361	20.821	1.00	30.44	C
ATOM	1324	C	TRP	A	179	38.328	20.385	21.789	1.00	30.36	C
ATOM	1325	O	TRP	A	179	39.008	19.446	21.416	1.00	29.60	O
ATOM	1326	CB	TRP	A	179	36.200	21.115	20.806	1.00	30.43	C
ATOM	1327	CG	TRP	A	179	35.426	22.183	20.136	1.00	29.86	C
ATOM	1328	CD1	TRP	A	179	35.882	23.409	19.721	1.00	27.42	C
ATOM	1329	CD2	TRP	A	179	34.054	22.115	19.769	1.00	29.51	C
ATOM	1330	NE1	TRP	A	179	34.870	24.102	19.104	1.00	30.05	N
ATOM	1331	CE2	TRP	A	179	33.728	23.338	19.130	1.00	30.70	C
ATOM	1332	CE3	TRP	A	179	33.067	21.145	19.902	1.00	27.24	C
ATOM	1333	CZ2	TRP	A	179	32.470	23.602	18.626	1.00	27.82	C
ATOM	1334	CZ3	TRP	A	179	31.829	21.397	19.389	1.00	28.43	C
ATOM	1335	CH2	TRP	A	179	31.532	22.630	18.759	1.00	29.15	C
ATOM	1336	N	GLY	A	180	38.087	20.634	23.065	1.00	30.98	N
ATOM	1337	CA	GLY	A	180	38.515	19.723	24.104	1.00	30.78	C
ATOM	1338	C	GLY	A	180	37.468	18.637	24.246	1.00	31.74	C
ATOM	1339	O	GLY	A	180	36.621	18.448	23.343	1.00	31.97	O
ATOM	1340	N	GLN	A	181	37.498	17.941	25.378	1.00	32.16	N
ATOM	1341	CA	GLN	A	181	36.628	16.798	25.608	1.00	33.12	C
ATOM	1342	C	GLN	A	181	35.161	17.108	25.885	1.00	32.52	C
ATOM	1343	O	GLN	A	181	34.813	18.181	26.348	1.00	32.55	O
ATOM	1344	CB	GLN	A	181	37.161	15.962	26.773	1.00	33.24	C
ATOM	1345	CG	GLN	A	181	36.780	16.496	28.155	1.00	37.52	C
ATOM	1346	CD	GLN	A	181	37.066	15.488	29.282	1.00	42.63	C
ATOM	1347	OE1	GLN	A	181	38.200	15.034	29.442	1.00	46.67	O
ATOM	1348	NE2	GLN	A	181	36.037	15.131	30.043	1.00	45.13	N
ATOM	1349	N	LEU	A	182	34.310	16.134	25.588	1.00	32.12	N
ATOM	1350	CA	LEU	A	182	32.907	16.155	26.001	1.00	31.88	C
ATOM	1351	C	LEU	A	182	32.928	16.114	27.528	1.00	30.66	C
ATOM	1352	O	LEU	A	182	33.481	15.190	28.087	1.00	30.31	O
ATOM	1353	CB	LEU	A	182	32.228	14.880	25.512	1.00	31.80	C
ATOM	1354	CG	LEU	A	182	30.715	14.723	25.454	1.00	33.59	C
ATOM	1355	CD1	LEU	A	182	30.319	13.250	25.572	1.00	32.15	C
ATOM	1356	CD2	LEU	A	182	30.059	15.454	26.520	1.00	36.60	C
ATOM	1357	N	THR	A	183	32.373	17.112	28.207	1.00	29.62	N
ATOM	1358	CA	THR	A	183	32.383	17.088	29.657	1.00	28.62	C
ATOM	1359	C	THR	A	183	31.088	16.509	30.123	1.00	28.48	C
ATOM	1360	O	THR	A	183	31.034	15.991	31.207	1.00	28.46	O
ATOM	1361	CB	THR	A	183	32.504	18.509	30.276	1.00	28.82	C
ATOM	1362	OG1	THR	A	183	31.441	19.364	29.795	1.00	28.93	O

ATOM	1363	CG2	THR	A	183	33.764	19.178	29.846	1.00	28.28
ATOM	1364	N	SER	A	184	30.005	16.673	29.355	1.00	28.14
ATOM	1365	CA	SER	A	184	28.734	16.144	29.805	1.00	28.16
ATOM	1366	C	SER	A	184	27.602	16.374	28.878	1.00	28.07
ATOM	1367	O	SER	A	184	27.703	17.120	27.931	1.00	29.72
ATOM	1368	CB	SER	A	184	28.357	16.750	31.149	1.00	27.81
ATOM	1369	OG	SER	A	184	28.166	18.132	31.017	1.00	30.35
ATOM	1370	N	ASN	A	185	26.505	15.721	29.176	1.00	27.93
ATOM	1371	CA	ASN	A	185	25.288	15.850	28.434	1.00	28.66
ATOM	1372	C	ASN	A	185	24.235	16.086	29.476	1.00	28.67
ATOM	1373	O	ASN	A	185	23.974	15.226	30.291	1.00	27.67
ATOM	1374	CB	ASN	A	185	24.927	14.543	27.683	1.00	28.53
ATOM	1375	CG	ASN	A	185	25.914	14.191	26.589	1.00	29.19
ATOM	1376	OD1	ASN	A	185	26.684	13.278	26.766	1.00	30.93
ATOM	1377	ND2	ASN	A	185	25.891	14.910	25.451	1.00	29.67
ATOM	1378	N	LEU	A	186	23.583	17.221	29.415	1.00	29.39
ATOM	1379	CA	LEU	A	186	22.557	17.526	30.388	1.00	30.47
ATOM	1380	C	LEU	A	186	21.179	17.177	29.835	1.00	30.85
ATOM	1381	O	LEU	A	186	20.796	17.566	28.714	1.00	30.64
ATOM	1382	CB	LEU	A	186	22.617	18.998	30.730	1.00	30.48
ATOM	1383	CG	LEU	A	186	21.779	19.484	31.917	1.00	32.51
ATOM	1384	CD1	LEU	A	186	22.330	18.952	33.238	1.00	31.55
ATOM	1385	CD2	LEU	A	186	21.678	21.072	31.954	1.00	29.96
ATOM	1386	N	LEU	A	187	20.409	16.439	30.613	1.00	31.21
ATOM	1387	CA	LEU	A	187	19.042	16.155	30.188	1.00	30.60
ATOM	1388	C	LEU	A	187	18.162	17.256	30.787	1.00	30.79
ATOM	1389	O	LEU	A	187	18.257	17.557	31.991	1.00	29.69
ATOM	1390	CB	LEU	A	187	18.626	14.781	30.644	1.00	30.50
ATOM	1391	CG	LEU	A	187	17.130	14.465	30.590	1.00	32.29
ATOM	1392	CD1	LEU	A	187	16.592	14.409	29.164	1.00	33.95
ATOM	1393	CD2	LEU	A	187	16.864	13.137	31.270	1.00	32.62
ATOM	1394	N	LEU	A	188	17.390	17.933	29.939	1.00	30.75
ATOM	1395	CA	LEU	A	188	16.513	18.973	30.426	1.00	31.42
ATOM	1396	C	LEU	A	188	15.053	18.646	30.107	1.00	32.02
ATOM	1397	O	LEU	A	188	14.676	18.481	28.941	1.00	31.44
ATOM	1398	CB	LEU	A	188	16.856	20.323	29.821	1.00	31.69
ATOM	1399	CG	LEU	A	188	18.253	20.823	30.131	1.00	32.24
ATOM	1400	CD1	LEU	A	188	19.094	20.883	28.904	1.00	32.79
ATOM	1401	CD2	LEU	A	188	18.121	22.200	30.658	1.00	32.69
ATOM	1402	N	ILE	A	189	14.228	18.568	31.146	1.00	32.26
ATOM	1403	CA	ILE	A	189	12.816	18.305	30.944	1.00	32.23
ATOM	1404	C	ILE	A	189	12.067	19.386	31.664	1.00	32.02
ATOM	1405	O	ILE	A	189	12.194	19.520	32.895	1.00	31.42
ATOM	1406	CB	ILE	A	189	12.427	16.966	31.479	1.00	32.05
ATOM	1407	CG1	ILE	A	189	13.256	15.894	30.793	1.00	33.02
ATOM	1408	CG2	ILE	A	189	10.957	16.764	31.232	1.00	32.90
ATOM	1409	CD1	ILE	A	189	13.006	14.466	31.277	1.00	34.20
ATOM	1410	N	GLY	A	190	11.279	20.145	30.898	1.00	31.94
ATOM	1411	CA	GLY	A	190	10.623	21.334	31.414	1.00	31.49
ATOM	1412	C	GLY	A	190	9.147	21.362	31.125	1.00	31.41
ATOM	1413	O	GLY	A	190	8.671	20.649	30.245	1.00	30.40
ATOM	1414	N	MET	A	191	8.433	22.154	31.914	1.00	31.55
ATOM	1415	CA	MET	A	191	7.027	22.378	31.697	1.00	32.41
ATOM	1416	C	MET	A	191	6.863	23.547	30.751	1.00	32.03
ATOM	1417	O	MET	A	191	7.722	24.435	30.658	1.00	32.29
ATOM	1418	CB	MET	A	191	6.313	22.672	33.008	1.00	33.21
ATOM	1419	CG	MET	A	191	6.294	21.499	33.957	1.00	35.09
ATOM	1420	SD	MET	A	191	5.828	21.953	35.652	1.00	39.51
ATOM	1421	CE	MET	A	191	4.213	22.512	35.350	1.00	39.52
ATOM	1422	N	GLU	A	192	5.755	23.539	30.033	1.00	31.96
ATOM	1423	CA	GLU	A	192	5.459	24.588	29.081	1.00	32.31

ATOM	1424	C	GLU	A	192	5.511	25.919	29.776	1.00	31.73
ATOM	1425	O	GLU	A	192	5.096	26.041	30.913	1.00	31.73
ATOM	1426	CB	GLU	A	192	4.087	24.375	28.508	1.00	33.09
ATOM	1427	CG	GLU	A	192	2.995	24.434	29.552	1.00	34.74
ATOM	1428	CD	GLU	A	192	1.679	23.940	29.019	1.00	36.73
ATOM	1429	OE1	GLU	A	192	1.630	23.515	27.832	1.00	37.30
ATOM	1430	OE2	GLU	A	192	0.698	24.001	29.791	1.00	38.64
ATOM	1431	N	GLY	A	193	6.049	26.926	29.103	1.00	31.79
ATOM	1432	CA	GLY	A	193	6.201	28.242	29.714	1.00	30.87
ATOM	1433	C	GLY	A	193	7.480	28.432	30.521	1.00	30.28
ATOM	1434	O	GLY	A	193	7.777	29.534	30.901	1.00	30.29
ATOM	1435	N	ASN	A	194	8.223	27.375	30.825	1.00	29.85
ATOM	1436	CA	ASN	A	194	9.460	27.536	31.604	1.00	29.38
ATOM	1437	C	ASN	A	194	10.473	28.371	30.840	1.00	28.61
ATOM	1438	O	ASN	A	194	10.606	28.216	29.615	1.00	28.58
ATOM	1439	CB	ASN	A	194	10.136	26.186	31.866	1.00	29.15
ATOM	1440	CG	ASN	A	194	9.565	25.445	33.049	1.00	29.50
ATOM	1441	OD1	ASN	A	194	8.632	25.896	33.714	1.00	32.61
ATOM	1442	ND2	ASN	A	194	10.146	24.305	33.334	1.00	28.53
ATOM	1443	N	VAL	A	195	11.221	29.194	31.561	1.00	27.70
ATOM	1444	CA	VAL	A	195	12.272	29.998	30.962	1.00	27.80
ATOM	1445	C	VAL	A	195	13.586	29.858	31.701	1.00	27.07
ATOM	1446	O	VAL	A	195	13.636	29.898	32.939	1.00	27.43
ATOM	1447	CB	VAL	A	195	11.904	31.520	31.018	1.00	28.18
ATOM	1448	CG1	VAL	A	195	13.056	32.371	30.560	1.00	26.78
ATOM	1449	CG2	VAL	A	195	10.662	31.789	30.187	1.00	29.06
ATOM	1450	N	THR	A	196	14.659	29.712	30.953	1.00	26.48
ATOM	1451	CA	THR	A	196	15.994	29.811	31.530	1.00	26.08
ATOM	1452	C	THR	A	196	16.462	31.184	31.084	1.00	27.03
ATOM	1453	O	THR	A	196	16.614	31.425	29.876	1.00	26.51
ATOM	1454	CB	THR	A	196	16.915	28.811	30.934	1.00	25.96
ATOM	1455	OG1	THR	A	196	16.525	27.478	31.309	1.00	26.20
ATOM	1456	CG2	THR	A	196	18.340	29.012	31.457	1.00	23.99
ATOM	1457	N	PRO	A	197	16.647	32.090	32.036	1.00	27.21
ATOM	1458	CA	PRO	A	197	17.053	33.451	31.725	1.00	27.46
ATOM	1459	C	PRO	A	197	18.432	33.512	31.113	1.00	27.48
ATOM	1460	O	PRO	A	197	19.248	32.588	31.293	1.00	27.52
ATOM	1461	CB	PRO	A	197	17.007	34.166	33.075	1.00	27.56
ATOM	1462	CG	PRO	A	197	16.226	33.344	33.924	1.00	28.15
ATOM	1463	CD	PRO	A	197	16.384	31.914	33.470	1.00	27.92
ATOM	1464	N	ALA	A	198	18.668	34.616	30.413	1.00	26.77
ATOM	1465	CA	ALA	A	198	19.877	34.830	29.647	1.00	26.92
ATOM	1466	C	ALA	A	198	21.172	34.681	30.438	1.00	27.14
ATOM	1467	O	ALA	A	198	21.354	35.278	31.520	1.00	25.53
ATOM	1468	CB	ALA	A	198	19.828	36.235	29.021	1.00	27.07
ATOM	1469	N	HIS	A	199	22.091	33.929	29.839	1.00	27.28
ATOM	1470	CA	HIS	A	199	23.399	33.673	30.424	1.00	27.81
ATOM	1471	C	HIS	A	199	24.319	33.208	29.333	1.00	28.09
ATOM	1472	O	HIS	A	199	23.857	32.933	28.197	1.00	28.34
ATOM	1473	CB	HIS	A	199	23.323	32.549	31.461	1.00	27.10
ATOM	1474	CG	HIS	A	199	22.963	31.221	30.864	1.00	29.70
ATOM	1475	ND1	HIS	A	199	21.668	30.891	30.525	1.00	30.01
ATOM	1476	CD2	HIS	A	199	23.728	30.180	30.463	1.00	29.60
ATOM	1477	CE1	HIS	A	199	21.647	29.684	29.993	1.00	29.66
ATOM	1478	NE2	HIS	A	199	22.881	29.236	29.926	1.00	30.43
ATOM	1479	N	TYR	A	200	25.612	33.104	29.671	1.00	28.30
ATOM	1480	CA	TYR	A	200	26.607	32.518	28.769	1.00	28.15
ATOM	1481	C	TYR	A	200	27.298	31.351	29.482	1.00	27.91
ATOM	1482	O	TYR	A	200	27.335	31.333	30.690	1.00	27.87
ATOM	1483	CB	TYR	A	200	27.585	33.534	28.217	1.00	28.02
ATOM	1484	CG	TYR	A	200	28.540	34.142	29.226	1.00	29.58

C
O
C
C
C
O
O
N
C
C
O
N
C
C
O
N
N
C
C
O
C
C
C
N
C
C
C
O
C
C
N
C
C
C
O
C
C
N
C
C
C
N
N
C
C
C
C
C

ATOM	1485	CD1	TYR	A	200	29.784	33.571	29.462	1.00	28.14	C
ATOM	1486	CD2	TYR	A	200	28.231	35.342	29.871	1.00	29.32	C
ATOM	1487	CE1	TYR	A	200	30.660	34.128	30.367	1.00	30.20	C
ATOM	1488	CE2	TYR	A	200	29.107	35.919	30.777	1.00	29.61	C
ATOM	1489	CZ	TYR	A	200	30.319	35.315	31.031	1.00	29.78	C
ATOM	1490	OH	TYR	A	200	31.180	35.864	31.940	1.00	24.43	O
ATOM	1491	N	ASP	A	201	27.797	30.363	28.727	1.00	27.73	N
ATOM	1492	CA	ASP	A	201	28.461	29.180	29.302	1.00	27.65	C
ATOM	1493	C	ASP	A	201	29.873	29.220	28.799	1.00	27.87	C
ATOM	1494	O	ASP	A	201	30.080	29.765	27.768	1.00	27.68	O
ATOM	1495	CB	ASP	A	201	27.775	27.884	28.855	1.00	26.32	C
ATOM	1496	CG	ASP	A	201	26.356	27.768	29.363	1.00	25.12	C
ATOM	1497	OD1	ASP	A	201	26.156	27.722	30.605	1.00	23.49	O
ATOM	1498	OD2	ASP	A	201	25.360	27.687	28.589	1.00	27.18	O
ATOM	1499	N	GLU	A	202	30.843	28.682	29.520	1.00	29.28	N
ATOM	1500	CA	GLU	A	202	32.228	28.672	29.013	1.00	31.18	C
ATOM	1501	C	GLU	A	202	32.571	27.435	28.190	1.00	31.54	C
ATOM	1502	O	GLU	A	202	33.734	27.060	28.139	1.00	33.48	O
ATOM	1503	CB	GLU	A	202	33.250	28.741	30.155	1.00	30.78	C
ATOM	1504	CG	GLU	A	202	33.122	29.981	31.003	1.00	33.61	C
ATOM	1505	CD	GLU	A	202	34.194	30.081	32.062	1.00	35.96	C
ATOM	1506	OE1	GLU	A	202	34.036	29.531	33.166	1.00	41.25	O
ATOM	1507	OE2	GLU	A	202	35.199	30.718	31.788	1.00	39.13	O
ATOM	1508	N	GLN	A	203	31.582	26.747	27.641	1.00	31.19	N
ATOM	1509	CA	GLN	A	203	31.844	25.589	26.833	1.00	30.78	C
ATOM	1510	C	GLN	A	203	31.078	25.743	25.556	1.00	29.96	C
ATOM	1511	O	GLN	A	203	30.213	26.581	25.462	1.00	30.22	O
ATOM	1512	CB	GLN	A	203	31.427	24.315	27.546	1.00	31.32	C
ATOM	1513	CG	GLN	A	203	32.364	23.971	28.725	1.00	34.97	C
ATOM	1514	CD	GLN	A	203	32.204	22.548	29.288	1.00	35.23	C
ATOM	1515	OE1	GLN	A	203	32.161	21.559	28.551	1.00	36.82	O
ATOM	1516	NE2	GLN	A	203	32.160	22.456	30.600	1.00	35.56	N
ATOM	1517	N	GLN	A	204	31.461	24.961	24.555	1.00	29.46	N
ATOM	1518	CA	GLN	A	204	30.791	24.911	23.292	1.00	28.65	C
ATOM	1519	C	GLN	A	204	29.641	23.941	23.494	1.00	28.27	C
ATOM	1520	O	GLN	A	204	29.797	22.941	24.194	1.00	27.65	O
ATOM	1521	CB	GLN	A	204	31.717	24.381	22.214	1.00	29.02	C
ATOM	1522	CG	GLN	A	204	33.006	25.154	22.028	1.00	28.50	C
ATOM	1523	CD	GLN	A	204	32.818	26.366	21.150	1.00	27.16	C
ATOM	1524	OE1	GLN	A	204	31.690	26.742	20.846	1.00	22.90	O
ATOM	1525	NE2	GLN	A	204	33.921	26.952	20.714	1.00	23.20	N
ATOM	1526	N	ASN	A	205	28.497	24.219	22.864	1.00	27.36	N
ATOM	1527	CA	ASN	A	205	27.302	23.424	23.058	1.00	27.13	C
ATOM	1528	C	ASN	A	205	26.552	23.052	21.777	1.00	27.39	C
ATOM	1529	O	ASN	A	205	26.151	23.930	20.999	1.00	27.36	O
ATOM	1530	CB	ASN	A	205	26.378	24.229	23.967	1.00	26.95	C
ATOM	1531	CG	ASN	A	205	25.083	23.525	24.327	1.00	27.30	C
ATOM	1532	OD1	ASN	A	205	24.726	22.453	23.818	1.00	27.11	O
ATOM	1533	ND2	ASN	A	205	24.319	24.190	25.171	1.00	24.64	N
ATOM	1534	N	PHE	A	206	26.383	21.749	21.542	1.00	26.46	N
ATOM	1535	CA	PHE	A	206	25.417	21.334	20.554	1.00	25.88	C
ATOM	1536	C	PHE	A	206	24.181	20.946	21.373	1.00	25.65	C
ATOM	1537	O	PHE	A	206	24.230	19.988	22.178	1.00	25.16	O
ATOM	1538	CB	PHE	A	206	25.909	20.170	19.717	1.00	26.28	C
ATOM	1539	CG	PHE	A	206	26.837	20.566	18.636	1.00	24.81	C
ATOM	1540	CD1	PHE	A	206	26.416	21.374	17.633	1.00	26.27	C
ATOM	1541	CD2	PHE	A	206	28.122	20.098	18.616	1.00	25.53	C
ATOM	1542	CE1	PHE	A	206	27.248	21.736	16.634	1.00	27.05	C
ATOM	1543	CE2	PHE	A	206	28.991	20.469	17.617	1.00	26.80	C
ATOM	1544	CZ	PHE	A	206	28.549	21.274	16.619	1.00	28.34	C
ATOM	1545	N	PHE	A	207	23.084	21.648	21.086	1.00	25.08	N

ATOM	1546	CA	PHE	A	207	21.809	21.625	21.827	1.00	25.52
ATOM	1547	C	PHE	A	207	20.775	20.885	21.011	1.00	25.55
ATOM	1548	O	PHE	A	207	20.261	21.410	20.058	1.00	25.61
ATOM	1549	CB	PHE	A	207	21.408	23.107	22.074	1.00	25.37
ATOM	1550	CG	PHE	A	207	20.146	23.346	22.872	1.00	23.70
ATOM	1551	CD1	PHE	A	207	18.938	23.547	22.234	1.00	24.08
ATOM	1552	CD2	PHE	A	207	20.199	23.551	24.220	1.00	25.24
ATOM	1553	CE1	PHE	A	207	17.800	23.864	22.927	1.00	24.50
ATOM	1554	CE2	PHE	A	207	19.035	23.883	24.959	1.00	26.04
ATOM	1555	CZ	PHE	A	207	17.836	24.021	24.298	1.00	26.68
ATOM	1556	N	ALA	A	208	20.490	19.657	21.422	1.00	26.60
ATOM	1557	CA	ALA	A	208	19.667	18.710	20.689	1.00	26.45
ATOM	1558	C	ALA	A	208	18.231	18.594	21.210	1.00	27.10
ATOM	1559	O	ALA	A	208	17.966	17.987	22.273	1.00	26.47
ATOM	1560	CB	ALA	A	208	20.303	17.363	20.766	1.00	25.67
ATOM	1561	N	GLN	A	209	17.306	19.121	20.419	1.00	27.02
ATOM	1562	CA	GLN	A	209	15.918	19.125	20.833	1.00	27.37
ATOM	1563	C	GLN	A	209	15.276	17.781	20.519	1.00	27.55
ATOM	1564	O	GLN	A	209	15.489	17.190	19.427	1.00	26.51
ATOM	1565	CB	GLN	A	209	15.195	20.301	20.179	1.00	27.15
ATOM	1566	CG	GLN	A	209	13.806	20.508	20.662	1.00	27.77
ATOM	1567	CD	GLN	A	209	13.740	20.959	22.126	1.00	29.40
ATOM	1568	OE1	GLN	A	209	14.773	21.166	22.774	1.00	27.27
ATOM	1569	NE2	GLN	A	209	12.517	21.118	22.636	1.00	27.15
ATOM	1570	N	ILE	A	210	14.461	17.327	21.473	1.00	28.32
ATOM	1571	CA	ILE	A	210	13.897	15.998	21.429	1.00	29.37
ATOM	1572	C	ILE	A	210	12.403	15.966	21.435	1.00	30.37
ATOM	1573	O	ILE	A	210	11.849	15.275	20.619	1.00	31.58
ATOM	1574	CB	ILE	A	210	14.413	15.204	22.605	1.00	29.80
ATOM	1575	CG1	ILE	A	210	15.830	14.734	22.302	1.00	30.27
ATOM	1576	CG2	ILE	A	210	13.525	14.005	22.864	1.00	30.25
ATOM	1577	CD1	ILE	A	210	16.624	14.421	23.516	1.00	32.28
ATOM	1578	N	LYS	A	211	11.757	16.664	22.374	1.00	31.18
ATOM	1579	CA	LYS	A	211	10.300	16.723	22.438	1.00	31.16
ATOM	1580	C	LYS	A	211	9.887	18.137	22.706	1.00	31.19
ATOM	1581	O	LYS	A	211	10.495	18.800	23.523	1.00	31.21
ATOM	1582	CB	LYS	A	211	9.767	15.891	23.591	1.00	32.00
ATOM	1583	CG	LYS	A	211	8.240	15.758	23.629	1.00	32.36
ATOM	1584	CD	LYS	A	211	7.787	15.177	24.970	1.00	33.23
ATOM	1585	CE	LYS	A	211	6.497	14.368	24.883	1.00	35.14
ATOM	1586	NZ	LYS	A	211	5.506	14.755	23.835	1.00	35.43
ATOM	1587	N	GLY	A	212	8.844	18.592	22.018	1.00	31.20
ATOM	1588	CA	GLY	A	212	8.337	19.934	22.174	1.00	30.96
ATOM	1589	C	GLY	A	212	9.209	20.975	21.495	1.00	31.42
ATOM	1590	O	GLY	A	212	10.167	20.670	20.771	1.00	31.06
ATOM	1591	N	TYR	A	213	8.857	22.224	21.734	1.00	31.43
ATOM	1592	CA	TYR	A	213	9.507	23.319	21.057	1.00	31.80
ATOM	1593	C	TYR	A	213	10.046	24.332	22.043	1.00	31.00
ATOM	1594	O	TYR	A	213	9.411	24.649	23.038	1.00	29.31
ATOM	1595	CB	TYR	A	213	8.502	23.959	20.127	1.00	32.64
ATOM	1596	CG	TYR	A	213	8.103	23.039	19.015	1.00	35.60
ATOM	1597	CD1	TYR	A	213	7.089	22.071	19.174	1.00	38.37
ATOM	1598	CD2	TYR	A	213	8.758	23.110	17.813	1.00	37.73
ATOM	1599	CE1	TYR	A	213	6.765	21.223	18.137	1.00	39.37
ATOM	1600	CE2	TYR	A	213	8.443	22.292	16.792	1.00	39.75
ATOM	1601	CZ	TYR	A	213	7.460	21.364	16.924	1.00	41.66
ATOM	1602	OH	TYR	A	213	7.232	20.603	15.782	1.00	46.53
ATOM	1603	N	LYS	A	214	11.260	24.787	21.777	1.00	30.99
ATOM	1604	CA	LYS	A	214	11.886	25.804	22.594	1.00	31.15
ATOM	1605	C	LYS	A	214	12.305	26.993	21.762	1.00	31.00
ATOM	1606	O	LYS	A	214	12.914	26.838	20.695	1.00	31.84

ATOM	1607	CB	LYS	A	214	13.112	25.244	23.311	1.00	31.08	
ATOM	1608	CG	LYS	A	214	12.806	24.432	24.557	1.00	30.96	C
ATOM	1609	CD	LYS	A	214	14.106	23.949	25.159	1.00	30.84	C
ATOM	1610	CE	LYS	A	214	13.982	23.555	26.584	1.00	29.59	C
ATOM	1611	NZ	LYS	A	214	15.278	23.118	27.162	1.00	28.16	N
ATOM	1612	N	ARG	A	215	11.992	28.193	22.241	1.00	30.59	N
ATOM	1613	CA	ARG	A	215	12.445	29.385	21.549	1.00	29.98	C
ATOM	1614	C	ARG	A	215	13.764	29.790	22.154	1.00	29.65	C
ATOM	1615	O	ARG	A	215	13.850	29.982	23.349	1.00	29.61	O
ATOM	1616	CB	ARG	A	215	11.431	30.502	21.690	1.00	30.08	C
ATOM	1617	CG	ARG	A	215	11.835	31.810	21.004	1.00	29.82	C
ATOM	1618	CD	ARG	A	215	11.221	32.960	21.731	1.00	32.78	C
ATOM	1619	NE	ARG	A	215	11.211	34.189	20.975	1.00	33.81	N
ATOM	1620	CZ	ARG	A	215	10.540	35.259	21.336	1.00	35.12	C
ATOM	1621	NH1	ARG	A	215	9.828	35.262	22.441	1.00	35.28	N
ATOM	1622	NH2	ARG	A	215	10.586	36.338	20.581	1.00	39.28	N
ATOM	1623	N	CYS	A	216	14.791	29.909	21.327	1.00	29.86	N
ATOM	1624	CA	CYS	A	216	16.129	30.237	21.797	1.00	29.64	C
ATOM	1625	C	CYS	A	216	16.533	31.603	21.274	1.00	30.06	C
ATOM	1626	O	CYS	A	216	16.516	31.807	20.092	1.00	31.32	O
ATOM	1627	CB	CYS	A	216	17.113	29.183	21.275	1.00	29.78	C
ATOM	1628	SG	CYS	A	216	16.658	27.438	21.641	1.00	30.35	S
ATOM	1629	N	ILE	A	217	16.849	32.556	22.149	1.00	30.28	N
ATOM	1630	CA	ILE	A	217	17.303	33.876	21.744	1.00	29.78	C
ATOM	1631	C	ILE	A	217	18.758	34.017	22.203	1.00	29.42	C
ATOM	1632	O	ILE	A	217	19.050	33.852	23.385	1.00	28.77	O
ATOM	1633	CB	ILE	A	217	16.427	34.969	22.391	1.00	30.51	C
ATOM	1634	CG1	ILE	A	217	14.934	34.705	22.111	1.00	31.29	C
ATOM	1635	CG2	ILE	A	217	16.785	36.313	21.843	1.00	30.11	C
ATOM	1636	CD1	ILE	A	217	14.009	35.655	22.847	1.00	33.08	C
ATOM	1637	N	LEU	A	218	19.647	34.326	21.257	1.00	28.60	N
ATOM	1638	CA	LEU	A	218	21.059	34.392	21.501	1.00	28.59	C
ATOM	1639	C	LEU	A	218	21.606	35.781	21.257	1.00	28.74	C
ATOM	1640	O	LEU	A	218	21.102	36.522	20.412	1.00	29.64	O
ATOM	1641	CB	LEU	A	218	21.788	33.422	20.574	1.00	28.52	C
ATOM	1642	CG	LEU	A	218	21.927	31.980	21.028	1.00	28.46	C
ATOM	1643	CD1	LEU	A	218	20.569	31.318	21.233	1.00	28.95	C
ATOM	1644	CD2	LEU	A	218	22.707	31.196	20.026	1.00	27.94	C
ATOM	1645	N	PHE	A	219	22.659	36.145	21.986	1.00	28.61	N
ATOM	1646	CA	PHE	A	219	23.309	37.458	21.774	1.00	28.07	C
ATOM	1647	C	PHE	A	219	24.811	37.221	21.710	1.00	28.25	C
ATOM	1648	O	PHE	A	219	25.352	36.468	22.517	1.00	28.82	O
ATOM	1649	CB	PHE	A	219	22.987	38.415	22.908	1.00	26.73	C
ATOM	1650	CG	PHE	A	219	21.522	38.553	23.199	1.00	27.28	C
ATOM	1651	CD1	PHE	A	219	20.874	37.658	24.029	1.00	25.73	C
ATOM	1652	CD2	PHE	A	219	20.782	39.598	22.648	1.00	28.57	C
ATOM	1653	CE1	PHE	A	219	19.536	37.811	24.311	1.00	26.59	C
ATOM	1654	CE2	PHE	A	219	19.443	39.722	22.898	1.00	27.78	C
ATOM	1655	CZ	PHE	A	219	18.816	38.828	23.725	1.00	28.22	C
ATOM	1656	N	PRO	A	220	25.501	37.840	20.776	1.00	28.30	N
ATOM	1657	CA	PRO	A	220	26.946	37.641	20.675	1.00	28.76	C
ATOM	1658	C	PRO	A	220	27.688	38.195	21.881	1.00	29.07	C
ATOM	1659	O	PRO	A	220	27.172	39.019	22.661	1.00	29.11	O
ATOM	1660	CB	PRO	A	220	27.336	38.413	19.426	1.00	28.79	C
ATOM	1661	CG	PRO	A	220	26.016	38.847	18.795	1.00	29.65	C
ATOM	1662	CD	PRO	A	220	24.987	38.794	19.790	1.00	28.85	C
ATOM	1663	N	PRO	A	221	28.914	37.727	22.057	1.00	29.09	N
ATOM	1664	CA	PRO	A	221	29.725	38.147	23.188	1.00	28.85	C
ATOM	1665	C	PRO	A	221	29.979	39.656	23.199	1.00	28.60	C
ATOM	1666	O	PRO	A	221	30.223	40.182	24.270	1.00	26.85	O
ATOM	1667	CB	PRO	A	221	31.018	37.412	22.975	1.00	29.01	C

ATOM	1668	CG	PRO	A	221	30.665	36.286	22.114	1.00	30.46
ATOM	1669	CD	PRO	A	221	29.580	36.722	21.232	1.00	29.52
ATOM	1670	N	ASP	A	222	29.853	40.335	22.062	1.00	28.17
ATOM	1671	CA	ASP	A	222	30.155	41.763	22.044	1.00	28.79
ATOM	1672	C	ASP	A	222	28.976	42.554	22.533	1.00	28.84
ATOM	1673	O	ASP	A	222	28.948	43.767	22.432	1.00	30.46
ATOM	1674	CB	ASP	A	222	30.631	42.287	20.693	1.00	27.27
ATOM	1675	CG	ASP	A	222	29.541	42.288	19.655	1.00	30.76
ATOM	1676	OD1	ASP	A	222	28.368	41.926	19.920	1.00	30.56
ATOM	1677	OD2	ASP	A	222	29.785	42.609	18.495	1.00	39.36
ATOM	1678	N	GLN	A	223	28.000	41.885	23.083	1.00	29.33
ATOM	1679	CA	GLN	A	223	26.902	42.614	23.676	1.00	29.61
ATOM	1680	C	GLN	A	223	27.012	42.533	25.194	1.00	28.97
ATOM	1681	O	GLN	A	223	26.065	42.825	25.919	1.00	29.82
ATOM	1682	CB	GLN	A	223	25.575	42.131	23.110	1.00	30.22
ATOM	1683	CG	GLN	A	223	25.244	42.854	21.762	1.00	33.01
ATOM	1684	CD	GLN	A	223	23.866	42.562	21.241	1.00	37.71
ATOM	1685	OE1	GLN	A	223	22.899	42.573	22.005	1.00	43.11
ATOM	1686	NE2	GLN	A	223	23.760	42.285	19.943	1.00	38.91
ATOM	1687	N	PHE	A	224	28.181	42.148	25.672	1.00	27.69
ATOM	1688	CA	PHE	A	224	28.452	42.168	27.115	1.00	28.30
ATOM	1689	C	PHE	A	224	27.932	43.497	27.771	1.00	29.27
ATOM	1690	O	PHE	A	224	27.248	43.448	28.786	1.00	29.24
ATOM	1691	CB	PHE	A	224	29.968	42.025	27.373	1.00	26.64
ATOM	1692	CG	PHE	A	224	30.338	41.881	28.810	1.00	27.05
ATOM	1693	CD1	PHE	A	224	30.526	42.992	29.619	1.00	26.91
ATOM	1694	CD2	PHE	A	224	30.560	40.649	29.365	1.00	25.77
ATOM	1695	CE1	PHE	A	224	30.878	42.864	30.930	1.00	24.95
ATOM	1696	CE2	PHE	A	224	30.928	40.522	30.675	1.00	25.92
ATOM	1697	CZ	PHE	A	224	31.056	41.641	31.467	1.00	27.45
ATOM	1698	N	GLU	A	225	28.253	44.658	27.177	1.00	30.23
ATOM	1699	CA	GLU	A	225	27.873	45.951	27.753	1.00	31.35
ATOM	1700	C	GLU	A	225	26.362	46.120	27.909	1.00	30.48
ATOM	1701	O	GLU	A	225	25.925	46.920	28.738	1.00	28.49
ATOM	1702	CB	GLU	A	225	28.401	47.130	26.927	1.00	32.35
ATOM	1703	CG	GLU	A	225	29.894	47.356	27.083	1.00	37.72
ATOM	1704	CD	GLU	A	225	30.327	48.801	27.406	1.00	45.22
ATOM	1705	OE1	GLU	A	225	29.848	49.472	28.389	1.00	44.87
ATOM	1706	OE2	GLU	A	225	31.230	49.248	26.665	1.00	54.04
ATOM	1707	N	CYS	A	226	25.579	45.375	27.116	1.00	29.44
ATOM	1708	CA	CYS	A	226	24.132	45.532	27.155	1.00	28.60
ATOM	1709	C	CYS	A	226	23.429	44.524	28.056	1.00	28.86

[illegible]

ATOM	1729	CE1	TYR	A	228	24.826	47.828	32.248	1.00	27.44
ATOM	1730	CE2	TYR	A	228	23.160	47.018	30.745	1.00	27.63
ATOM	1731	CZ	TYR	A	228	23.964	48.040	31.186	1.00	29.01
ATOM	1732	OH	TYR	A	228	23.924	49.260	30.536	1.00	28.98
ATOM	1733	N	PRO	A	229	26.009	41.101	33.261	1.00	28.11
ATOM	1734	CA	PRO	A	229	26.146	39.900	34.082	1.00	28.54
ATOM	1735	C	PRO	A	229	26.424	40.223	35.534	1.00	27.88
ATOM	1736	O	PRO	A	229	27.085	41.212	35.786	1.00	28.01
ATOM	1737	CB	PRO	A	229	27.396	39.214	33.494	1.00	28.67
ATOM	1738	CG	PRO	A	229	28.116	40.314	32.802	1.00	28.96
ATOM	1739	CD	PRO	A	229	27.046	41.191	32.231	1.00	28.25
ATOM	1740	N	TYR	A	230	25.981	39.372	36.453	1.00	26.80
ATOM	1741	CA	TYR	A	230	26.296	39.542	37.840	1.00	25.91
ATOM	1742	C	TYR	A	230	27.838	39.508	38.028	1.00	25.72
ATOM	1743	O	TYR	A	230	28.582	39.094	37.167	1.00	25.79
ATOM	1744	CB	TYR	A	230	25.673	38.431	38.677	1.00	25.13
ATOM	1745	CG	TYR	A	230	24.193	38.523	38.900	1.00	24.95
ATOM	1746	CD1	TYR	A	230	23.289	38.036	37.933	1.00	24.25
ATOM	1747	CD2	TYR	A	230	23.675	39.063	40.090	1.00	25.16
ATOM	1748	CE1	TYR	A	230	21.944	38.105	38.139	1.00	24.40
ATOM	1749	CE2	TYR	A	230	22.311	39.114	40.320	1.00	23.86
ATOM	1750	CZ	TYR	A	230	21.464	38.647	39.336	1.00	26.74
ATOM	1751	OH	TYR	A	230	20.122	38.713	39.501	1.00	34.04
ATOM	1752	N	PRO	A	231	28.309	39.966	39.167	1.00	25.05
ATOM	1753	CA	PRO	A	231	29.711	39.806	39.517	1.00	24.73
ATOM	1754	C	PRO	A	231	30.151	38.318	39.500	1.00	24.94
ATOM	1755	O	PRO	A	231	29.359	37.410	39.817	1.00	24.30
ATOM	1756	CB	PRO	A	231	29.760	40.353	40.957	1.00	25.28
ATOM	1757	CG	PRO	A	231	28.613	41.331	41.016	1.00	25.28
ATOM	1758	CD	PRO	A	231	27.535	40.687	40.198	1.00	24.73
ATOM	1759	N	VAL	A	232	31.419	38.086	39.173	1.00	23.75
ATOM	1760	CA	VAL	A	232	31.898	36.766	39.055	1.00	24.45
ATOM	1761	C	VAL	A	232	31.725	35.935	40.351	1.00	24.83
ATOM	1762	O	VAL	A	232	31.402	34.754	40.263	1.00	24.83
ATOM	1763	CB	VAL	A	232	33.375	36.761	38.582	1.00	25.13
ATOM	1764	CG1	VAL	A	232	33.974	35.405	38.828	1.00	23.91
ATOM	1765	CG2	VAL	A	232	33.481	37.091	37.092	1.00	23.86
ATOM	1766	N	HIS	A	233	31.886	36.540	41.525	1.00	25.24
ATOM	1767	CA	HIS	A	233	31.766	35.806	42.790	1.00	26.43
ATOM	1768	C	HIS	A	233	30.321	35.656	43.304	1.00	26.47
ATOM	1769	O	HIS	A	233	30.052	34.986	44.289	1.00	26.08
ATOM	1770	CB	HIS	A	233	32.617	36.475	43.855	1.00	26.65
ATOM	1771	CG	HIS	A	233	34.078	36.428	43.564	1.00	28.51
ATOM	1772	ND1	HIS	A	233	34.764	37.493	43.013	1.00	30.95
ATOM	1773	CD2	HIS	A	233	34.987	35.435	43.730	1.00	29.75
ATOM	1774	CE1	HIS	A	233	36.043	37.169	42.888	1.00	31.18
ATOM	1775	NE2	HIS	A	233	36.201	35.919	43.302	1.00	32.10
ATOM	1776	N	HIS	A	234	29.391	36.283	42.626	1.00	26.94
ATOM	1777	CA	HIS	A	234	27.999	36.141	42.983	1.00	27.89
ATOM	1778	C	HIS	A	234	27.456	34.838	42.411	1.00	27.85
ATOM	1779	O	HIS	A	234	27.871	34.403	41.384	1.00	27.14
ATOM	1780	CB	HIS	A	234	27.219	37.258	42.372	1.00	28.20
ATOM	1781	CG	HIS	A	234	25.782	37.279	42.748	1.00	27.95
ATOM	1782	ND1	HIS	A	234	24.840	36.514	42.092	1.00	30.05
ATOM	1783	CD2	HIS	A	234	25.105	38.057	43.624	1.00	26.43
ATOM	1784	CE1	HIS	A	234	23.644	36.789	42.588	1.00	29.87
ATOM	1785	NE2	HIS	A	234	23.781	37.714	43.524	1.00	27.85
ATOM	1786	N	PRO	A	235	26.561	34.197	43.130	1.00	28.44
ATOM	1787	CA	PRO	A	235	26.018	32.929	42.678	1.00	28.87
ATOM	1788	C	PRO	A	235	25.431	32.956	41.294	1.00	28.34
ATOM	1789	O	PRO	A	235	25.387	31.884	40.716	1.00	28.04

ATOM	1790	CB	PRO A 235	24.931	32.636	43.713	1.00	29.59
ATOM	1791	CG	PRO A 235	25.473	33.290	44.975	1.00	29.10
ATOM	1792	CD	PRO A 235	26.059	34.571	44.466	1.00	28.17
ATOM	1793	N	CYS A 236	25.021	34.112	40.770	1.00	27.29
ATOM	1794	CA	CYS A 236	24.447	34.135	39.455	1.00	26.15
ATOM	1795	C	CYS A 236	25.478	34.602	38.451	1.00	25.39
ATOM	1796	O	CYS A 236	25.140	35.102	37.361	1.00	26.21
ATOM	1797	CB	CYS A 236	23.158	34.923	39.436	1.00	26.36
ATOM	1798	SG	CYS A 236	21.908	34.207	40.555	1.00	29.77
ATOM	1799	N	ASP A 237	26.746	34.411	38.803	1.00	24.07
ATOM	1800	CA	ASP A 237	27.830	34.528	37.855	1.00	24.53
ATOM	1801	C	ASP A 237	27.441	33.977	36.467	1.00	24.31
ATOM	1802	O	ASP A 237	26.966	32.865	36.344	1.00	23.00
ATOM	1803	CB	ASP A 237	29.018	33.769	38.353	1.00	24.42
ATOM	1804	CG	ASP A 237	30.233	33.850	37.414	1.00	28.30
ATOM	1805	OD1	ASP A 237	30.433	34.877	36.673	1.00	29.51
ATOM	1806	OD2	ASP A 237	31.071	32.900	37.386	1.00	29.51
ATOM	1807	N	ARG A 238	27.633	34.803	35.445	1.00	24.85
ATOM	1808	CA	ARG A 238	27.388	34.434	34.041	1.00	26.59
ATOM	1809	C	ARG A 238	25.921	34.663	33.595	1.00	25.95
ATOM	1810	O	ARG A 238	25.647	34.628	32.426	1.00	26.55
ATOM	1811	CB	ARG A 238	27.834	32.994	33.737	1.00	26.57
ATOM	1812	CG	ARG A 238	29.311	32.758	33.891	1.00	27.77
ATOM	1813	CD	ARG A 238	29.727	31.360	33.399	1.00	28.78
ATOM	1814	NE	ARG A 238	29.142	30.434	34.337	1.00	33.68
ATOM	1815	CZ	ARG A 238	28.046	29.736	34.140	1.00	36.98
ATOM	1816	NH1	ARG A 238	27.383	29.782	32.980	1.00	38.06
ATOM	1817	NH2	ARG A 238	27.609	28.987	35.129	1.00	37.87
ATOM	1818	N	GLN A 239	25.012	34.933	34.516	1.00	25.97
ATOM	1819	CA	GLN A 239	23.634	35.245	34.152	1.00	26.05
ATOM	1820	C	GLN A 239	23.478	36.776	34.037	1.00	26.67
ATOM	1821	O	GLN A 239	24.196	37.531	34.699	1.00	27.33
ATOM	1822	CB	GLN A 239	22.663	34.724	35.201	1.00	25.55
ATOM	1823	CG	GLN A 239	23.027	33.417	35.843	1.00	27.73
ATOM	1824	CD	GLN A 239	23.246	32.282	34.859	1.00	31.42
ATOM	1825	OE1	GLN A 239	22.332	31.881	34.133	1.00	34.43
ATOM	1826	NE2	GLN A 239	24.457	31.761	34.835	1.00	31.35
ATOM	1827	N	SER A 240	22.569	37.233	33.185	1.00	26.94
ATOM	1828	CA	SER A 240	22.293	38.653	33.034	1.00	26.93
ATOM	1829	C	SER A 240	21.490	39.118	34.229	1.00	26.78
ATOM	1830	O	SER A 240	20.642	38.389	34.703	1.00	27.00
ATOM	1831	CB	SER A 240	21.414	38.891	31.818	1.00	26.95
ATOM	1832	OG	SER A 240	20.916	40.234	31.804	1.00	27.75
ATOM	1833	N	GLN A 241	21.733	40.331	34.696	1.00	26.90
ATOM	1834	CA	GLN A 241	20.963	40.903	35.816	1.00	27.25
ATOM	1835	C	GLN A 241	19.632	41.520	35.311	1.00	27.84
ATOM	1836	O	GLN A 241	18.763	41.866	36.104	1.00	27.28
ATOM	1837	CB	GLN A 241	21.734	42.009	36.546	1.00	26.21
ATOM	1838	CG	GLN A 241	22.971	41.636	37.344	1.00	26.53
ATOM	1839	CD	GLN A 241	23.740	42.877	37.817	1.00	26.65
ATOM	1840	OE1	GLN A 241	23.436	43.425	38.859	1.00	31.02
ATOM	1841	NE2	GLN A 241	24.714	43.318	37.048	1.00	28.00
ATOM	1842	N	VAL A 242	19.464	41.657	34.004	1.00	28.19
ATOM	1843	CA	VAL A 242	18.284	42.335	33.552	1.00	28.55
ATOM	1844	C	VAL A 242	17.117	41.409	33.563	1.00	28.88
ATOM	1845	O	VAL A 242	17.198	40.314	33.053	1.00	28.92
ATOM	1846	CB	VAL A 242	18.422	42.770	32.093	1.00	29.31
ATOM	1847	CG1	VAL A 242	17.165	43.549	31.647	1.00	27.90
ATOM	1848	CG2	VAL A 242	19.711	43.530	31.866	1.00	28.67
ATOM	1849	N	ASP A 243	16.009	41.862	34.113	1.00	28.55
ATOM	1850	CA	ASP A 243	14.766	41.103	34.052	1.00	27.62

C
C
C
N
C
C
C
O
C
S
N
C
C
O
C
C
O
O
O
N
C
C
O
C
C
N
C
N
N
N
C
C
O
C
C
C
O
N
C
C
O
C
C
O
N
N
C
C
O
C
C
C
O
N
N
C
C
O
C
C
C
N
C

ATOM	1912	CA	GLU	A	250	17.994	53.202	34.789	1.00	35.66
ATOM	1913	C	GLU	A	250	19.249	52.494	35.235	1.00	34.69
ATOM	1914	O	GLU	A	250	20.313	53.077	35.207	1.00	34.29
ATOM	1915	CB	GLU	A	250	17.252	53.685	36.021	1.00	36.82
ATOM	1916	CG	GLU	A	250	16.115	54.609	35.650	1.00	41.99
ATOM	1917	CD	GLU	A	250	15.554	55.360	36.827	1.00	48.02
ATOM	1918	OE1	GLU	A	250	16.114	55.275	37.949	1.00	53.95
ATOM	1919	OE2	GLU	A	250	14.543	56.043	36.614	1.00	52.47
ATOM	1920	N	ARG	A	251	19.139	51.254	35.691	1.00	33.87
ATOM	1921	CA	ARG	A	251	20.335	50.550	36.088	1.00	33.26
ATOM	1922	C	ARG	A	251	21.013	49.992	34.873	1.00	32.36
ATOM	1923	O	ARG	A	251	22.210	49.914	34.849	1.00	32.79
ATOM	1924	CB	ARG	A	251	20.040	49.425	37.066	1.00	34.36
ATOM	1925	CG	ARG	A	251	19.476	49.895	38.425	1.00	36.13
ATOM	1926	CD	ARG	A	251	18.966	48.771	39.339	1.00	39.31
ATOM	1927	NE	ARG	A	251	20.052	47.860	39.733	1.00	42.09
ATOM	1928	CZ	ARG	A	251	19.890	46.563	39.964	1.00	41.90
ATOM	1929	NH1	ARG	A	251	18.696	46.006	39.843	1.00	41.62
ATOM	1930	NH2	ARG	A	251	20.932	45.815	40.296	1.00	41.98
ATOM	1931	N	PHE	A	252	20.272	49.654	33.825	1.00	31.66
ATOM	1932	CA	PHE	A	252	20.879	48.977	32.677	1.00	30.28
ATOM	1933	C	PHE	A	252	20.528	49.651	31.347	1.00	29.72
ATOM	1934	O	PHE	A	252	19.882	49.096	30.483	1.00	29.84
ATOM	1935	CB	PHE	A	252	20.420	47.531	32.675	1.00	29.99
ATOM	1936	CG	PHE	A	252	20.405	46.864	34.050	1.00	29.63
ATOM	1937	CD1	PHE	A	252	21.562	46.717	34.796	1.00	29.74
ATOM	1938	CD2	PHE	A	252	19.238	46.317	34.554	1.00	31.30
ATOM	1939	CE1	PHE	A	252	21.546	46.067	36.039	1.00	30.44
ATOM	1940	CE2	PHE	A	252	19.199	45.668	35.805	1.00	31.42
ATOM	1941	CZ	PHE	A	252	20.356	45.551	36.550	1.00	32.04
ATOM	1942	N	PRO	A	253	21.034	50.844	31.155	1.00	29.72
ATOM	1943	CA	PRO	A	253	20.607	51.662	30.022	1.00	28.89
ATOM	1944	C	PRO	A	253	20.953	50.996	28.689	1.00	29.46
ATOM	1945	O	PRO	A	253	20.166	51.061	27.722	1.00	28.81
ATOM	1946	CB	PRO	A	253	21.361	52.975	30.240	1.00	28.37
ATOM	1947	CG	PRO	A	253	22.543	52.614	31.155	1.00	27.97
ATOM	1948	CD	PRO	A	253	22.136	51.463	31.942	1.00	29.24
ATOM	1949	N	ASN	A	254	22.093	50.323	28.590	1.00	29.37
ATOM	1950	CA	ASN	A	254	22.401	49.742	27.274	1.00	28.85
ATOM	1951	C	ASN	A	254	21.547	48.529	26.902	1.00	28.28
ATOM	1952	O	ASN	A	254	21.663	47.970	25.794	1.00	27.47
ATOM	1953	CB	ASN	A	254	23.874	49.407	27.122	1.00	28.84
ATOM	1954	CG	ASN	A	254	24.745	50.654	26.996	1.00	29.96
ATOM	1955	OD1	ASN	A	254	25.622	50.882	27.834	1.00	31.21
ATOM	1956	ND2	ASN	A	254	24.531	51.443	25.931	1.00	29.17
ATOM	1957	N	PHE	A	255	20.687	48.091	27.797	1.00	27.30
ATOM	1958	CA	PHE	A	255	19.801	47.006	27.368	1.00	27.53
ATOM	1959	C	PHE	A	255	18.844	47.512	26.283	1.00	26.80
ATOM	1960	O	PHE	A	255	18.193	46.750	25.603	1.00	27.22
ATOM	1961	CB	PHE	A	255	19.015	46.448	28.538	1.00	27.48
ATOM	1962	CG	PHE	A	255	18.282	45.250	28.198	1.00	26.63
ATOM	1963	CD1	PHE	A	255	18.950	44.073	27.998	1.00	29.74
ATOM	1964	CD2	PHE	A	255	16.935	45.288	28.039	1.00	26.29
ATOM	1965	CE1	PHE	A	255	18.273	42.947	27.681	1.00	29.50
ATOM	1966	CE2	PHE	A	255	16.261	44.192	27.692	1.00	26.21
ATOM	1967	CZ	PHE	A	255	16.917	43.012	27.518	1.00	30.10
ATOM	1968	N	GLN	A	256	18.776	48.818	26.125	1.00	26.57
ATOM	1969	CA	GLN	A	256	17.918	49.429	25.137	1.00	26.55
ATOM	1970	C	GLN	A	256	18.522	49.253	23.745	1.00	26.46
ATOM	1971	O	GLN	A	256	17.881	49.560	22.771	1.00	24.39
ATOM	1972	CB	GLN	A	256	17.756	50.933	25.423	1.00	26.18

[illegible]

ATOM	1973	CG	GLN	A	256	16.656	51.226	26.406	1.00	28.72
ATOM	1974	CD	GLN	A	256	16.721	52.636	27.028	1.00	29.64
ATOM	1975	OE1	GLN	A	256	15.737	53.358	27.005	1.00	32.56
ATOM	1976	NE2	GLN	A	256	17.861	52.999	27.590	1.00	26.88
ATOM	1977	N	ASN	A	257	19.774	48.809	23.694	1.00	27.86
ATOM	1978	CA	ASN	A	257	20.509	48.611	22.452	1.00	29.22
ATOM	1979	C	ASN	A	257	20.733	47.143	22.116	1.00	30.46
ATOM	1980	O	ASN	A	257	21.399	46.855	21.132	1.00	32.28
ATOM	1981	CB	ASN	A	257	21.897	49.232	22.510	1.00	28.50
ATOM	1982	CG	ASN	A	257	21.878	50.666	22.890	1.00	29.30
ATOM	1983	OD1	ASN	A	257	22.329	51.025	23.989	1.00	27.33
ATOM	1984	ND2	ASN	A	257	21.382	51.523	21.982	1.00	25.68
ATOM	1985	N	VAL	A	258	20.200	46.213	22.900	1.00	31.51
ATOM	1986	CA	VAL	A	258	20.432	44.791	22.623	1.00	32.54
ATOM	1987	C	VAL	A	258	19.674	44.295	21.386	1.00	32.39
ATOM	1988	O	VAL	A	258	18.543	44.766	21.081	1.00	32.29
ATOM	1989	CB	VAL	A	258	20.050	43.937	23.841	1.00	33.12
ATOM	1990	CG1	VAL	A	258	18.546	43.818	23.986	1.00	32.40
ATOM	1991	CG2	VAL	A	258	20.646	42.566	23.686	1.00	36.78
ATOM	1992	N	VAL	A	259	20.302	43.386	20.654	1.00	32.45
ATOM	1993	CA	VAL	A	259	19.682	42.796	19.448	1.00	32.96
ATOM	1994	C	VAL	A	259	19.943	41.301	19.391	1.00	33.08
ATOM	1995	O	VAL	A	259	21.074	40.878	19.351	1.00	33.50
ATOM	1996	CB	VAL	A	259	20.269	43.401	18.172	1.00	33.19
ATOM	1997	CG1	VAL	A	259	19.625	42.771	16.962	1.00	34.73
ATOM	1998	CG2	VAL	A	259	20.054	44.901	18.143	1.00	32.25
ATOM	1999	N	GLY	A	260	18.902	40.491	19.406	1.00	33.70
ATOM	2000	CA	GLY	A	260	19.091	39.052	19.368	1.00	34.16
ATOM	2001	C	GLY	A	260	19.036	38.327	18.014	1.00	33.92
ATOM	2002	O	GLY	A	260	18.655	38.891	16.975	1.00	33.59
ATOM	2003	N	TYR	A	261	19.469	37.071	18.052	1.00	33.14
ATOM	2004	CA	TYR	A	261	19.375	36.157	16.938	1.00	33.55
ATOM	2005	C	TYR	A	261	18.439	35.095	17.489	1.00	33.41
ATOM	2006	O	TYR	A	261	18.700	34.542	18.561	1.00	33.67
ATOM	2007	CB	TYR	A	261	20.715	35.508	16.657	1.00	33.81
ATOM	2008	CG	TYR	A	261	21.758	36.416	16.061	1.00	36.80
ATOM	2009	CD1	TYR	A	261	21.886	36.545	14.694	1.00	42.37
ATOM	2010	CD2	TYR	A	261	22.602	37.139	16.851	1.00	38.31
ATOM	2011	CE1	TYR	A	261	22.823	37.378	14.139	1.00	43.02
ATOM	2012	CE2	TYR	A	261	23.556	37.938	16.312	1.00	40.85
ATOM	2013	CZ	TYR	A	261	23.660	38.058	14.955	1.00	42.50
ATOM	2014	OH	TYR	A	261	24.617	38.865	14.414	1.00	45.43
ATOM	2015	N	GLU	A	262	17.322	34.832	16.834	1.00	32.71
ATOM	2016	CA	GLU	A	262	16.394	33.886	17.438	1.00	32.21
ATOM	2017	C	GLU	A	262	15.975	32.768	16.508	1.00	30.89
ATOM	2018	O	GLU	A	262	16.180	32.819	15.294	1.00	30.63
ATOM	2019	CB	GLU	A	262	15.196	34.613	18.049	1.00	32.16
ATOM	2020	CG	GLU	A	262	13.968	34.753	17.210	1.00	30.84
ATOM	2021	CD	GLU	A	262	12.818	35.380	17.992	1.00	31.21
ATOM	2022	OE1	GLU	A	262	12.761	36.612	18.108	1.00	34.58
ATOM	2023	OE2	GLU	A	262	11.962	34.663	18.504	1.00	29.56
ATOM	2024	N	THR	A	263	15.447	31.746	17.133	1.00	28.97
ATOM	2025	CA	THR	A	263	14.941	30.587	16.447	1.00	28.44
ATOM	2026	C	THR	A	263	14.080	29.743	17.387	1.00	27.24
ATOM	2027	O	THR	A	263	14.088	29.893	18.620	1.00	26.43
ATOM	2028	CB	THR	A	263	16.093	29.746	15.902	1.00	28.50
ATOM	2029	OG1	THR	A	263	15.592	28.798	14.964	1.00	27.18
ATOM	2030	CG2	THR	A	263	16.697	28.854	16.975	1.00	28.63
ATOM	2031	N	VAL	A	264	13.306	28.887	16.776	1.00	27.22
ATOM	2032	CA	VAL	A	264	12.509	27.942	17.497	1.00	28.49
ATOM	2033	C	VAL	A	264	12.925	26.576	17.022	1.00	29.10

[illegible]

ATOM	2095	N	ILE	A	273	25.237	26.418	19.934	1.00	26.87	N
ATOM	2096	CA	ILE	A	273	25.381	27.591	20.810	1.00	26.89	C
ATOM	2097	C	ILE	A	273	26.841	27.751	21.182	1.00	27.37	C
ATOM	2098	O	ILE	A	273	27.359	27.049	22.056	1.00	27.62	O
ATOM	2099	CB	ILE	A	273	24.647	27.422	22.085	1.00	26.74	C
ATOM	2100	CG1	ILE	A	273	23.182	27.185	21.799	1.00	25.89	C
ATOM	2101	CG2	ILE	A	273	24.852	28.675	22.941	1.00	26.79	C
ATOM	2102	CD1	ILE	A	273	22.338	27.108	23.065	1.00	25.37	C
ATOM	2103	N	PRO	A	274	27.491	28.681	20.515	1.00	27.39	N
ATOM	2104	CA	PRO	A	274	28.923	28.871	20.601	1.00	27.76	C
ATOM	2105	C	PRO	A	274	29.287	29.415	21.934	1.00	27.42	C
ATOM	2106	O	PRO	A	274	28.483	30.135	22.518	1.00	26.71	O
ATOM	2107	CB	PRO	A	274	29.250	29.840	19.426	1.00	27.60	C
ATOM	2108	CG	PRO	A	274	27.980	30.107	18.741	1.00	28.16	C
ATOM	2109	CD	PRO	A	274	26.876	29.504	19.497	1.00	28.21	C
ATOM	2110	N	MET	A	275	30.467	29.017	22.394	1.00	27.88	N
ATOM	2111	CA	MET	A	275	30.979	29.346	23.717	1.00	28.91	C
ATOM	2112	C	MET	A	275	30.974	30.849	23.919	1.00	28.27	C
ATOM	2113	O	MET	A	275	31.284	31.589	23.000	1.00	27.69	O
ATOM	2114	CB	MET	A	275	32.365	28.808	23.869	1.00	29.00	C
ATOM	2115	CG	MET	A	275	32.861	28.842	25.313	1.00	34.03	C
ATOM	2116	SD	MET	A	275	34.499	28.084	25.455	1.00	39.22	S
ATOM	2117	CE	MET	A	275	35.200	28.479	23.871	1.00	36.40	C
ATOM	2118	N	TYR	A	276	30.531	31.294	25.086	1.00	27.79	N
ATOM	2119	CA	TYR	A	276	30.472	32.746	25.389	1.00	29.16	C
ATOM	2120	C	TYR	A	276	29.267	33.465	24.802	1.00	27.80	C
ATOM	2121	O	TYR	A	276	29.076	34.609	25.107	1.00	28.35	O
ATOM	2122	CB	TYR	A	276	31.779	33.506	24.974	1.00	29.63	C
ATOM	2123	CG	TYR	A	276	32.918	33.207	25.897	1.00	34.37	C
ATOM	2124	CD1	TYR	A	276	32.966	33.783	27.146	1.00	38.52	C
ATOM	2125	CD2	TYR	A	276	33.934	32.306	25.547	1.00	39.45	C
ATOM	2126	CE1	TYR	A	276	33.955	33.493	28.022	1.00	39.83	C
ATOM	2127	CE2	TYR	A	276	34.965	32.015	26.449	1.00	42.05	C
ATOM	2128	CZ	TYR	A	276	34.954	32.609	27.685	1.00	42.46	C
ATOM	2129	OH	TYR	A	276	35.949	32.343	28.624	1.00	48.60	O
ATOM	2130	N	TRP	A	277	28.468	32.830	23.951	1.00	27.45	N
ATOM	2131	CA	TRP	A	277	27.271	33.485	23.419	1.00	26.19	C
ATOM	2132	C	TRP	A	277	26.160	33.442	24.424	1.00	25.93	C
ATOM	2133	O	TRP	A	277	25.882	32.409	24.973	1.00	26.84	O
ATOM	2134	CB	TRP	A	277	26.796	32.826	22.130	1.00	26.07	C
ATOM	2135	CG	TRP	A	277	27.525	33.276	20.957	1.00	24.33	C
ATOM	2136	CD1	TRP	A	277	28.862	33.118	20.731	1.00	24.81	C
ATOM	2137	CD2	TRP	A	277	26.995	33.935	19.821	1.00	23.88	C
ATOM	2138	NE1	TRP	A	277	29.204	33.674	19.531	1.00	24.86	N
ATOM	2139	CE2	TRP	A	277	28.073	34.182	18.944	1.00	24.16	C
ATOM	2140	CE3	TRP	A	277	25.727	34.398	19.464	1.00	25.16	C
ATOM	2141	CZ2	TRP	A	277	27.923	34.858	17.747	1.00	23.53	C
ATOM	2142	CZ3	TRP	A	277	25.579	35.064	18.250	1.00	25.92	C
ATOM	2143	CH2	TRP	A	277	26.679	35.292	17.414	1.00	24.81	C
ATOM	2144	N	TRP	A	278	25.516	34.566	24.673	1.00	26.34	N
ATOM	2145	CA	TRP	A	278	24.408	34.623	25.625	1.00	26.96	C
ATOM	2146	C	TRP	A	278	23.229	33.847	25.050	1.00	27.33	C
ATOM	2147	O	TRP	A	278	23.021	33.846	23.847	1.00	27.18	O
ATOM	2148	CB	TRP	A	278	23.952	36.062	25.838	1.00	26.87	C
ATOM	2149	CG	TRP	A	278	24.965	36.916	26.455	1.00	27.61	C
ATOM	2150	CD1	TRP	A	278	25.997	37.558	25.833	1.00	28.27	C
ATOM	2151	CD2	TRP	A	278	25.065	37.228	27.830	1.00	25.52	C
ATOM	2152	NE1	TRP	A	278	26.731	38.261	26.753	1.00	28.77	N
ATOM	2153	CE2	TRP	A	278	26.164	38.072	27.991	1.00	29.02	C
ATOM	2154	CE3	TRP	A	278	24.303	36.915	28.942	1.00	26.86	C
ATOM	2155	CZ2	TRP	A	278	26.541	38.563	29.224	1.00	28.96	C

ATOM	2156	CH3	TRP	A	278	24.676	37.402	30.157	1.00	29.86
ATOM	2157	CH2	TRP	A	278	25.780	38.224	30.293	1.00	27.89
ATOM	2158	N	HIS	A	279	22.455	33.193	25.890	1.00	27.47
ATOM	2159	CA	HIS	A	279	21.263	32.554	25.380	1.00	28.50
ATOM	2160	C	HIS	A	279	20.158	32.574	26.403	1.00	28.09
ATOM	2161	O	HIS	A	279	20.420	32.445	27.584	1.00	29.26
ATOM	2162	CB	HIS	A	279	21.528	31.112	24.899	1.00	28.46
ATOM	2163	CG	HIS	A	279	22.339	30.278	25.838	1.00	30.29
ATOM	2164	ND1	HIS	A	279	23.712	30.381	25.928	1.00	31.17
ATOM	2165	CD2	HIS	A	279	21.984	29.270	26.673	1.00	31.89
ATOM	2166	CE1	HIS	A	279	24.161	29.484	26.790	1.00	31.48
ATOM	2167	NE2	HIS	A	279	23.136	28.810	27.271	1.00	30.92
ATOM	2168	N	HIS	A	280	18.942	32.764	25.916	1.00	28.12
ATOM	2169	CA	HIS	A	280	17.713	32.827	26.695	1.00	28.50
ATOM	2170	C	HIS	A	280	16.884	31.731	26.071	1.00	28.78
ATOM	2171	O	HIS	A	280	16.739	31.709	24.851	1.00	29.67
ATOM	2172	CB	HIS	A	280	17.069	34.196	26.498	1.00	27.71
ATOM	2173	CG	HIS	A	280	15.600	34.223	26.727	1.00	29.69
ATOM	2174	ND1	HIS	A	280	15.021	34.905	27.782	1.00	31.66
ATOM	2175	CD2	HIS	A	280	14.579	33.648	26.047	1.00	31.45
ATOM	2176	CE1	HIS	A	280	13.710	34.743	27.738	1.00	30.66
ATOM	2177	NE2	HIS	A	280	13.415	33.985	26.696	1.00	31.28
ATOM	2178	N	ILE	A	281	16.346	30.814	26.860	1.00	29.23
ATOM	2179	CA	ILE	A	281	15.651	29.652	26.294	1.00	29.56
ATOM	2180	C	ILE	A	281	14.311	29.497	26.922	1.00	30.16
ATOM	2181	O	ILE	A	281	14.190	29.517	28.135	1.00	30.14
ATOM	2182	CB	ILE	A	281	16.478	28.409	26.516	1.00	29.69
ATOM	2183	CG1	ILE	A	281	17.707	28.491	25.628	1.00	30.69
ATOM	2184	CG2	ILE	A	281	15.686	27.119	26.172	1.00	28.22
ATOM	2185	CD1	ILE	A	281	18.689	27.534	25.980	1.00	33.11
ATOM	2186	N	GLU	A	282	13.280	29.380	26.103	1.00	31.22
ATOM	2187	CA	GLU	A	282	11.931	29.278	26.665	1.00	31.99
ATOM	2188	C	GLU	A	282	11.123	28.182	26.007	1.00	31.79
ATOM	2189	O	GLU	A	282	11.131	28.021	24.806	1.00	32.77
ATOM	2190	CB	GLU	A	282	11.200	30.638	26.618	1.00	32.16
ATOM	2191	CG	GLU	A	282	11.017	31.265	25.253	1.00	33.49
ATOM	2192	CD	GLU	A	282	10.378	32.663	25.313	1.00	36.22
ATOM	2193	OE1	GLU	A	282	10.879	33.552	26.043	1.00	38.20
ATOM	2194	OE2	GLU	A	282	9.380	32.898	24.603	1.00	38.15
ATOM	2195	N	SER	A	283	10.454	27.392	26.827	1.00	31.81
ATOM	2196	CA	SER	A	283	9.592	26.333	26.331	1.00	30.79
ATOM	2197	C	SER	A	283	8.245	26.975	25.950	1.00	30.98
ATOM	2198	O	SER	A	283	7.635	27.640	26.780	1.00	30.36
ATOM	2199	CB	SER	A	283	9.430	25.305	27.424	1.00	30.62
ATOM	2200	OG	SER	A	283	10.586	24.463	27.496	1.00	29.45
ATOM	2201	N	LEU	A	284	7.776	26.790	24.709	1.00	31.06
ATOM	2202	CA	LEU	A	284	6.587	27.504	24.270	1.00	31.43
ATOM	2203	C	LEU	A	284	5.398	27.390	25.204	1.00	31.82
ATOM	2204	O	LEU	A	284	5.137	26.333	25.776	1.00	31.15
ATOM	2205	CB	LEU	A	284	6.156	27.098	22.895	1.00	31.75
ATOM	2206	CG	LEU	A	284	7.223	27.187	21.829	1.00	33.11
ATOM	2207	CD1	LEU	A	284	6.571	27.388	20.478	1.00	33.62
ATOM	2208	CD2	LEU	A	284	8.170	28.270	22.142	1.00	32.91
ATOM	2209	N	LEU	A	285	4.701	28.512	25.355	1.00	32.74
ATOM	2210	CA	LEU	A	285	3.481	28.564	26.147	1.00	34.35
ATOM	2211	C	LEU	A	285	2.502	27.591	25.540	1.00	34.74
ATOM	2212	O	LEU	A	285	2.375	27.510	24.332	1.00	34.70
ATOM	2213	CB	LEU	A	285	2.863	29.946	26.094	1.00	34.26
ATOM	2214	CG	LEU	A	285	3.729	31.060	26.638	1.00	34.74
ATOM	2215	CD1	LEU	A	285	3.183	32.369	26.160	1.00	34.14
ATOM	2216	CD2	LEU	A	285	3.780	30.970	28.136	1.00	34.97

ATOM	2217	N	ASN A 286	1.838	26.829	26.380	1.00	35.89
ATOM	2218	CA	ASN A 286	0.841	25.863	25.896	1.00	37.12
ATOM	2219	C	ASN A 286	1.363	24.809	24.909	1.00	36.14
ATOM	2220	O	ASN A 286	0.600	24.306	24.108	1.00	35.85
ATOM	2221	CB	ASN A 286	-0.336	26.635	25.274	1.00	37.80
ATOM	2222	CG	ASN A 286	-0.882	27.690	26.219	1.00	42.02
ATOM	2223	OD1	ASN A 286	-1.326	27.365	27.326	1.00	47.86
ATOM	2224	ND2	ASN A 286	-0.809	28.965	25.817	1.00	46.67
ATOM	2225	N	GLY A 287	2.653	24.478	24.974	1.00	35.43
ATOM	2226	CA	GLY A 287	3.262	23.539	24.057	1.00	34.11
ATOM	2227	C	GLY A 287	3.569	22.202	24.666	1.00	34.02
ATOM	2228	O	GLY A 287	4.073	21.317	23.977	1.00	34.98
ATOM	2229	N	GLY A 288	3.242	22.023	25.944	1.00	33.71
ATOM	2230	CA	GLY A 288	3.476	20.759	26.616	1.00	33.00
ATOM	2231	C	GLY A 288	4.908	20.692	27.107	1.00	33.19
ATOM	2232	O	GLY A 288	5.625	21.681	26.989	1.00	33.40
ATOM	2233	N	ILE A 289	5.337	19.540	27.621	1.00	32.87
ATOM	2234	CA	ILE A 289	6.647	19.425	28.195	1.00	33.57
ATOM	2235	C	ILE A 289	7.695	19.434	27.118	1.00	33.13
ATOM	2236	O	ILE A 289	7.454	19.023	25.995	1.00	34.41
ATOM	2237	CB	ILE A 289	6.809	18.137	29.006	1.00	34.25
ATOM	2238	CG1	ILE A 289	6.789	16.920	28.102	1.00	36.60
ATOM	2239	CG2	ILE A 289	5.746	18.005	30.113	1.00	34.91
ATOM	2240	CD1	ILE A 289	7.271	15.659	28.834	1.00	38.16
ATOM	2241	N	THR A 290	8.891	19.868	27.465	1.00	31.68
ATOM	2242	CA	THR A 290	9.956	19.853	26.498	1.00	29.79
ATOM	2243	C	THR A 290	11.054	18.938	26.958	1.00	27.97
ATOM	2244	O	THR A 290	11.256	18.739	28.133	1.00	26.58
ATOM	2245	CB	THR A 290	10.526	21.278	26.305	1.00	30.77
ATOM	2246	OG1	THR A 290	10.957	21.827	27.554	1.00	27.89
ATOM	2247	CG2	THR A 290	9.458	22.243	25.789	1.00	30.82
ATOM	2248	N	ILE A 291	11.787	18.395	26.021	1.00	26.96
ATOM	2249	CA	ILE A 291	12.910	17.594	26.383	1.00	27.83
ATOM	2250	C	ILE A 291	14.059	17.952	25.493	1.00	27.31
ATOM	2251	O	ILE A 291	13.880	18.028	24.289	1.00	27.83
ATOM	2252	CB	ILE A 291	12.596	16.104	26.195	1.00	28.72
ATOM	2253	CG1	ILE A 291	11.480	15.671	27.140	1.00	29.09
ATOM	2254	CG2	ILE A 291	13.865	15.278	26.438	1.00	28.55
ATOM	2255	CD1	ILE A 291	11.115	14.232	26.986	1.00	30.86
ATOM	2256	N	THR A 292	15.240	18.093	26.074	1.00	26.63
ATOM	2257	CA	THR A 292	16.415	18.444	25.333	1.00	26.88
ATOM	2258	C	THR A 292	17.587	17.724	25.915	1.00	26.52
ATOM	2259	O	THR A 292	17.631	17.524	27.102	1.00	26.40
ATOM	2260	CB	THR A 292	16.753	19.971	25.532	1.00	27.41
ATOM	2261	OG1	THR A 292	15.652	20.841	25.187	1.00	28.64
ATOM	2262	CG2	THR A 292	17.852	20.398	24.592	1.00	28.03
ATOM	2263	N	VAL A 293	18.577	17.414	25.093	1.00	26.32
ATOM	2264	CA	VAL A 293	19.834	16.896	25.585	1.00	26.86
ATOM	2265	C	VAL A 293	20.964	17.718	24.978	1.00	27.24
ATOM	2266	O	VAL A 293	21.011	17.892	23.757	1.00	26.60
ATOM	2267	CB	VAL A 293	20.029	15.422	25.226	1.00	27.56
ATOM	2268	CG1	VAL A 293	21.472	15.004	25.411	1.00	27.42
ATOM	2269	CG2	VAL A 293	19.138	14.546	26.126	1.00	29.64
ATOM	2270	N	ASN A 294	21.849	18.278	25.814	1.00	27.26
ATOM	2271	CA	ASN A 294	22.960	19.055	25.258	1.00	27.81
ATOM	2272	C	ASN A 294	24.253	18.268	25.281	1.00	27.79
ATOM	2273	O	ASN A 294	24.338	17.177	25.883	1.00	28.63
ATOM	2274	CB	ASN A 294	23.118	20.446	25.881	1.00	27.69
ATOM	2275	CG	ASN A 294	23.703	20.405	27.287	1.00	30.08
ATOM	2276	OD1	ASN A 294	24.309	19.397	27.677	1.00	29.80
ATOM	2277	ND2	ASN A 294	23.513	21.506	28.064	1.00	28.15

ATOM	2278	N	PHE	A	295	25.235	18.816	24.576	1.00	27.46
ATOM	2279	CA	PHE	A	295	26.548	18.219	24.428	1.00	27.53
ATOM	2280	C	PHE	A	295	27.543	19.352	24.675	1.00	28.01
ATOM	2281	O	PHE	A	295	27.720	20.239	23.811	1.00	28.48
ATOM	2282	CB	PHE	A	295	26.756	17.703	23.009	1.00	27.33
ATOM	2283	CG	PHE	A	295	25.955	16.487	22.658	1.00	27.87
ATOM	2284	CD1	PHE	A	295	24.586	16.560	22.496	1.00	27.75
ATOM	2285	CD2	PHE	A	295	26.581	15.287	22.426	1.00	26.57
ATOM	2286	CE1	PHE	A	295	23.892	15.466	22.147	1.00	28.45
ATOM	2287	CE2	PHE	A	295	25.869	14.187	22.095	1.00	27.40
ATOM	2288	CZ	PHE	A	295	24.541	14.262	21.952	1.00	27.57
ATOM	2289	N	TRP	A	296	28.166	19.346	25.852	1.00	28.46
ATOM	2290	CA	TRP	A	296	29.097	20.402	26.249	1.00	29.05
ATOM	2291	C	TRP	A	296	30.545	19.950	26.077	1.00	29.39
ATOM	2292	O	TRP	A	296	30.981	18.942	26.663	1.00	29.90
ATOM	2293	CB	TRP	A	296	28.850	20.825	27.696	1.00	29.23
ATOM	2294	CG	TRP	A	296	27.917	21.982	27.908	1.00	31.21
ATOM	2295	CD1	TRP	A	296	27.805	23.080	27.135	1.00	33.45
ATOM	2296	CD2	TRP	A	296	26.995	22.171	28.996	1.00	33.70
ATOM	2297	NE1	TRP	A	296	26.864	23.941	27.651	1.00	33.91
ATOM	2298	CE2	TRP	A	296	26.344	23.403	28.787	1.00	33.49
ATOM	2299	CE3	TRP	A	296	26.638	21.414	30.113	1.00	36.30
ATOM	2300	CZ2	TRP	A	296	25.377	23.902	29.642	1.00	35.59
ATOM	2301	CZ3	TRP	A	296	25.643	21.899	30.967	1.00	37.80
ATOM	2302	CH2	TRP	A	296	25.034	23.141	30.727	1.00	37.39
ATOM	2303	N	TYR	A	297	31.288	20.720	25.292	1.00	29.87
ATOM	2304	CA	TYR	A	297	32.693	20.472	25.017	1.00	30.43
ATOM	2305	C	TYR	A	297	33.548	21.629	25.518	1.00	30.98
ATOM	2306	O	TYR	A	297	33.167	22.765	25.387	1.00	29.50
ATOM	2307	CB	TYR	A	297	32.909	20.334	23.502	1.00	30.36
ATOM	2308	CG	TYR	A	297	32.304	19.083	22.919	1.00	30.07
ATOM	2309	CD1	TYR	A	297	30.946	19.012	22.627	1.00	29.74
ATOM	2310	CD2	TYR	A	297	33.083	17.960	22.687	1.00	28.56
ATOM	2311	CE1	TYR	A	297	30.397	17.859	22.105	1.00	28.46
ATOM	2312	CE2	TYR	A	297	32.546	16.800	22.213	1.00	27.77
ATOM	2313	CZ	TYR	A	297	31.202	16.758	21.913	1.00	28.10
ATOM	2314	OH	TYR	A	297	30.654	15.613	21.438	1.00	27.72
ATOM	2315	N	LYS	A	298	34.723	21.325	26.060	1.00	32.77
ATOM	2316	CA	LYS	A	298	35.671	22.349	26.470	1.00	34.43
ATOM	2317	C	LYS	A	298	36.126	23.014	25.216	1.00	34.91
ATOM	2318	O	LYS	A	298	36.236	22.373	24.194	1.00	35.28
ATOM	2319	CB	LYS	A	298	36.865	21.749	27.221	1.00	35.25
ATOM	2320	CG	LYS	A	298	36.595	21.446	28.725	1.00	38.48
ATOM	2321	CD	LYS	A	298	37.834	20.990	29.532	1.00	42.40
ATOM	2322	CE	LYS	A	298	37.429	20.600	30.991	1.00	45.34
ATOM	2323	NZ	LYS	A	298	38.515	19.883	31.770	1.00	49.07
ATOM	2324	N	GLY	A	299	36.384	24.305	25.267	1.00	36.67
ATOM	2325	CA	GLY	A	299	36.837	24.998	24.080	1.00	38.01
ATOM	2326	C	GLY	A	299	38.249	24.618	23.689	1.00	39.33
ATOM	2327	O	GLY	A	299	38.965	23.987	24.437	1.00	39.19
ATOM	2328	N	ALA	A	300	38.644	25.017	22.498	1.00	41.68
ATOM	2329	CA	ALA	A	300	40.014	24.844	22.048	1.00	43.93
ATOM	2330	C	ALA	A	300	40.964	25.618	22.940	1.00	45.88
ATOM	2331	O	ALA	A	300	40.558	26.437	23.741	1.00	46.12
ATOM	2332	CB	ALA	A	300	40.148	25.343	20.639	1.00	43.86
ATOM	2333	N	PRO	A	301	42.254	25.393	22.773	1.00	49.19
ATOM	2334	CA	PRO	A	301	43.251	26.084	23.605	1.00	50.79
ATOM	2335	C	PRO	A	301	43.465	27.525	23.204	1.00	52.01
ATOM	2336	O	PRO	A	301	43.302	27.869	22.041	1.00	52.73
ATOM	2337	CB	PRO	A	301	44.534	25.325	23.303	1.00	50.51
ATOM	2338	CG	PRO	A	301	44.357	24.908	21.884	1.00	50.75

[illegible]

ATOM	2400	N	HIS	A	313	35.148	37.753	22.916	1.00	29.85
ATOM	2401	CA	HIS	A	313	35.125	36.727	23.931	1.00	30.47
ATOM	2402	C	HIS	A	313	36.410	36.710	24.689	1.00	29.99
ATOM	2403	O	HIS	A	313	36.439	36.320	25.846	1.00	29.25
ATOM	2404	CB	HIS	A	313	34.787	35.331	23.392	1.00	31.64
ATOM	2405	CG	HIS	A	313	35.860	34.714	22.566	1.00	35.54
ATOM	2406	ND1	HIS	A	313	35.987	34.950	21.208	1.00	41.10
ATOM	2407	CD2	HIS	A	313	36.844	33.846	22.891	1.00	40.77
ATOM	2408	CE1	HIS	A	313	37.047	34.303	20.750	1.00	40.02
ATOM	2409	NE2	HIS	A	313	37.567	33.603	21.743	1.00	41.82
ATOM	2410	N	GLN	A	314	37.482	37.137	24.056	1.00	29.68
ATOM	2411	CA	GLN	A	314	38.754	37.185	24.751	1.00	29.77
ATOM	2412	C	GLN	A	314	38.729	38.294	25.817	1.00	30.57
ATOM	2413	O	GLN	A	314	39.222	38.120	26.956	1.00	29.69
ATOM	2414	CB	GLN	A	314	39.893	37.401	23.743	1.00	30.02
ATOM	2415	CG	GLN	A	314	40.111	36.182	22.786	1.00	29.84
ATOM	2416	CD	GLN	A	314	41.126	36.437	21.687	1.00	31.76
ATOM	2417	OE1	GLN	A	314	40.982	37.377	20.885	1.00	32.81
ATOM	2418	NE2	GLN	A	314	42.160	35.594	21.637	1.00	34.23
ATOM	2419	N	LYS	A	315	38.111	39.421	25.471	1.00	30.44
ATOM	2420	CA	LYS	A	315	38.026	40.491	26.431	1.00	31.27
ATOM	2421	C	LYS	A	315	37.192	40.023	27.635	1.00	30.75
ATOM	2422	O	LYS	A	315	37.511	40.373	28.784	1.00	29.60
ATOM	2423	CB	LYS	A	315	37.461	41.752	25.808	1.00	31.76
ATOM	2424	CG	LYS	A	315	38.483	42.491	24.982	1.00	33.90
ATOM	2425	CD	LYS	A	315	37.902	43.711	24.329	1.00	37.97
ATOM	2426	CE	LYS	A	315	38.911	44.380	23.419	1.00	41.69
ATOM	2427	NZ	LYS	A	315	38.264	45.411	22.536	1.00	44.93
ATOM	2428	N	VAL	A	316	36.163	39.216	27.362	1.00	29.26
ATOM	2429	CA	VAL	A	316	35.318	38.731	28.422	1.00	29.23
ATOM	2430	C	VAL	A	316	36.152	37.841	29.339	1.00	29.87
ATOM	2431	O	VAL	A	316	36.075	37.926	30.584	1.00	30.33
ATOM	2432	CB	VAL	A	316	34.079	37.973	27.900	1.00	28.68
ATOM	2433	CG1	VAL	A	316	33.306	37.376	29.037	1.00	28.32
ATOM	2434	CG2	VAL	A	316	33.144	38.887	27.136	1.00	28.57
ATOM	2435	N	ALA	A	317	36.976	37.007	28.728	1.00	29.36
ATOM	2436	CA	ALA	A	317	37.878	36.136	29.487	1.00	29.26
ATOM	2437	C	ALA	A	317	38.776	36.941	30.389	1.00	28.08
ATOM	2438	O	ALA	A	317	39.052	36.564	31.519	1.00	27.65
ATOM	2439	CB	ALA	A	317	38.733	35.267	28.561	1.00	28.70
ATOM	2440	N	ILE	A	318	39.211	38.068	29.873	1.00	28.33
ATOM	2441	CA	ILE	A	318	40.083	38.945	30.628	1.00	28.40
ATOM	2442	C	ILE	A	318	39.338	39.526	31.825	1.00	28.94
ATOM	2443	O	ILE	A	318	39.880	39.503	32.924	1.00	29.70
ATOM	2444	CB	ILE	A	318	40.697	40.053	29.732	1.00	27.74
ATOM	2445	CG1	ILE	A	318	41.683	39.443	28.743	1.00	28.14
ATOM	2446	CG2	ILE	A	318	41.433	41.082	30.577	1.00	27.98
ATOM	2447	CD1	ILE	A	318	42.271	40.447	27.759	1.00	29.17
ATOM	2448	N	MET	A	319	38.115	40.032	31.630	1.00	28.75
ATOM	2449	CA	MET	A	319	37.402	40.666	32.735	1.00	28.64
ATOM	2450	C	MET	A	319	37.140	39.669	33.877	1.00	29.22
ATOM	2451	O	MET	A	319	37.308	39.981	35.082	1.00	30.32
ATOM	2452	CB	MET	A	319	36.123	41.355	32.273	1.00	28.33
ATOM	2453	CG	MET	A	319	36.327	42.553	31.334	1.00	27.88
ATOM	2454	SD	MET	A	319	34.779	43.349	30.857	1.00	29.90
ATOM	2455	CE	MET	A	319	34.146	42.288	29.455	1.00	28.71
ATOM	2456	N	ARG	A	320	36.802	38.446	33.515	1.00	29.22
ATOM	2457	CA	ARG	A	320	36.543	37.429	34.520	1.00	28.75
ATOM	2458	C	ARG	A	320	37.806	37.221	35.325	1.00	28.81
ATOM	2459	O	ARG	A	320	37.782	37.132	36.569	1.00	28.76
ATOM	2460	CB	ARG	A	320	36.142	36.118	33.861	1.00	28.67

N
C
C
O
C
C
N
C
C
N
N
C
C
O
C
C
O
N
N
C
C
O
C
C
C
N
C
C
C
O
C
C
C
N
C
C
C
O
C
C
C
N
C
C
C
O
C
C
C
N
C
C
C
O
C

ATOM	2461	CG	ARG	A	320	34.810	36.125	33.120	1.00	29.62	C
ATOM	2462	CD	ARG	A	320	34.371	34.747	32.603	1.00	27.74	C
ATOM	2463	NE	ARG	A	320	34.193	33.843	33.716	1.00	24.81	N
ATOM	2464	CZ	ARG	A	320	33.161	33.887	34.526	1.00	27.65	C
ATOM	2465	NH1	ARG	A	320	32.180	34.752	34.330	1.00	29.97	N
ATOM	2466	NH2	ARG	A	320	33.088	33.064	35.542	1.00	27.59	N
ATOM	2467	N	ASN	A	321	38.933	37.167	34.627	1.00	28.60	N
ATOM	2468	CA	ASN	A	321	40.185	36.884	35.320	1.00	28.16	C
ATOM	2469	C	ASN	A	321	40.526	38.003	36.291	1.00	28.13	C
ATOM	2470	O	ASN	A	321	40.945	37.742	37.423	1.00	29.13	O
ATOM	2471	CB	ASN	A	321	41.318	36.615	34.319	1.00	28.09	C
ATOM	2472	CG	ASN	A	321	41.317	35.177	33.817	1.00	27.94	C
ATOM	2473	OD1	ASN	A	321	40.924	34.286	34.555	1.00	28.53	O
ATOM	2474	ND2	ASN	A	321	41.751	34.946	32.560	1.00	23.58	N
ATOM	2475	N	ILE	A	322	40.337	39.250	35.876	1.00	27.49	N
ATOM	2476	CA	ILE	A	322	40.659	40.350	36.749	1.00	27.70	C
ATOM	2477	C	ILE	A	322	39.792	40.209	38.017	1.00	27.39	C
ATOM	2478	O	ILE	A	322	40.284	40.315	39.101	1.00	26.48	O
ATOM	2479	CB	ILE	A	322	40.353	41.680	36.059	1.00	28.04	C
ATOM	2480	CG1	ILE	A	322	41.302	41.941	34.897	1.00	30.79	C
ATOM	2481	CG2	ILE	A	322	40.426	42.832	37.038	1.00	28.09	C
ATOM	2482	CD1	ILE	A	322	42.756	41.766	35.226	1.00	32.31	C
ATOM	2483	N	GLU	A	323	38.494	39.961	37.854	1.00	27.37	N
ATOM	2484	CA	GLU	A	323	37.618	39.868	38.985	1.00	27.48	C
ATOM	2485	C	GLU	A	323	38.060	38.784	39.881	1.00	27.52	C
ATOM	2486	O	GLU	A	323	38.096	38.956	41.079	1.00	26.90	O
ATOM	2487	CB	GLU	A	323	36.183	39.637	38.558	1.00	27.48	C
ATOM	2488	CG	GLU	A	323	35.592	40.849	37.910	1.00	27.65	C
ATOM	2489	CD	GLU	A	323	34.199	40.631	37.318	1.00	25.69	C
ATOM	2490	OE1	GLU	A	323	33.236	40.358	38.050	1.00	24.42	O
ATOM	2491	OE2	GLU	A	323	34.072	40.808	36.101	1.00	24.98	O
ATOM	2492	N	LYS	A	324	38.419	37.659	39.297	1.00	29.13	N
ATOM	2493	CA	LYS	A	324	38.845	36.508	40.097	1.00	30.35	C
ATOM	2494	C	LYS	A	324	40.099	36.825	40.912	1.00	31.16	C
ATOM	2495	O	LYS	A	324	40.123	36.547	42.111	1.00	32.01	O
ATOM	2496	CB	LYS	A	324	39.116	35.289	39.222	1.00	30.16	C
ATOM	2497	CG	LYS	A	324	37.887	34.625	38.648	1.00	30.63	C
ATOM	2498	CD	LYS	A	324	38.344	33.454	37.765	1.00	30.70	C
ATOM	2499	CE	LYS	A	324	37.204	32.713	37.038	1.00	30.66	C
ATOM	2500	NZ	LYS	A	324	37.728	31.451	36.325	1.00	27.19	N
ATOM	2501	N	MET	A	325	41.119	37.401	40.270	1.00	31.84	N
ATOM	2502	CA	MET	A	325	42.392	37.735	40.939	1.00	32.71	C
ATOM	2503	C	MET	A	325	42.216	38.743	42.078	1.00	32.44	C
ATOM	2504	O	MET	A	325	42.840	38.637	43.141	1.00	30.36	O
ATOM	2505	CB	MET	A	325	43.417	38.276	39.923	1.00	32.98	C
ATOM	2506	CG	MET	A	325	43.978	37.216	39.032	1.00	36.70	C
ATOM	2507	SD	MET	A	325	44.734	37.879	37.532	1.00	46.02	S
ATOM	2508	CE	MET	A	325	45.616	39.197	38.250	1.00	46.11	C
ATOM	2509	N	LEU	A	326	41.376	39.738	41.837	1.00	33.01	N
ATOM	2510	CA	LEU	A	326	41.116	40.756	42.852	1.00	34.17	C
ATOM	2511	C	LEU	A	326	40.490	40.151	44.083	1.00	33.83	C
ATOM	2512	O	LEU	A	326	40.871	40.464	45.188	1.00	33.22	O
ATOM	2513	CB	LEU	A	326	40.185	41.818	42.305	1.00	34.61	C
ATOM	2514	CG	LEU	A	326	40.740	43.194	41.974	1.00	37.32	C
ATOM	2515	CD1	LEU	A	326	42.251	43.297	41.924	1.00	39.32	C
ATOM	2516	CD2	LEU	A	326	40.158	43.607	40.643	1.00	39.20	C
ATOM	2517	N	GLY	A	327	39.534	39.262	43.861	1.00	34.50	N
ATOM	2518	CA	GLY	A	327	38.842	38.569	44.930	1.00	34.87	C
ATOM	2519	C	GLY	A	327	39.796	37.776	45.779	1.00	35.34	C
ATOM	2520	O	GLY	A	327	39.728	37.810	47.016	1.00	35.51	O
ATOM	2521	N	GLU	A	328	40.725	37.081	45.139	1.00	35.88	N

ATOM	2522	CA	GLU A 328	41.708	36.346	45.926	1.00	36.96
ATOM	2523	C	GLU A 328	42.668	37.302	46.614	1.00	35.46
ATOM	2524	O	GLU A 328	42.958	37.144	47.799	1.00	35.65
ATOM	2525	CB	GLU A 328	42.465	35.340	45.074	1.00	37.89
ATOM	2526	CG	GLU A 328	41.555	34.272	44.481	1.00	43.40
ATOM	2527	CD	GLU A 328	41.694	32.905	45.135	1.00	50.39
ATOM	2528	OE1	GLU A 328	42.006	32.861	46.361	1.00	53.07
ATOM	2529	OE2	GLU A 328	41.503	31.876	44.396	1.00	53.24
ATOM	2530	N	ALA A 329	43.110	38.331	45.906	1.00	34.23
ATOM	2531	CA	ALA A 329	44.130	39.208	46.469	1.00	33.82
ATOM	2532	C	ALA A 329	43.652	40.031	47.632	1.00	33.70
ATOM	2533	O	ALA A 329	44.384	40.279	48.547	1.00	33.79
ATOM	2534	CB	ALA A 329	44.711	40.079	45.428	1.00	33.36
ATOM	2535	N	LEU A 330	42.410	40.451	47.602	1.00	34.57
ATOM	2536	CA	LEU A 330	41.869	41.243	48.694	1.00	35.04
ATOM	2537	C	LEU A 330	41.351	40.397	49.837	1.00	35.29
ATOM	2538	O	LEU A 330	41.011	40.922	50.884	1.00	35.39
ATOM	2539	CB	LEU A 330	40.721	42.096	48.182	1.00	34.85
ATOM	2540	CG	LEU A 330	41.141	43.081	47.095	1.00	35.83
ATOM	2541	CD1	LEU A 330	39.907	43.564	46.304	1.00	37.05
ATOM	2542	CD2	LEU A 330	41.842	44.244	47.662	1.00	34.12
ATOM	2543	N	GLY A 331	41.223	39.098	49.619	1.00	36.20
ATOM	2544	CA	GLY A 331	40.751	38.207	50.659	1.00	36.87
ATOM	2545	C	GLY A 331	39.260	38.250	50.957	1.00	37.48
ATOM	2546	O	GLY A 331	38.807	37.527	51.846	1.00	38.53
ATOM	2547	N	ASN A 332	38.509	39.085	50.242	1.00	37.67
ATOM	2548	CA	ASN A 332	37.069	39.175	50.399	1.00	38.06
ATOM	2549	C	ASN A 332	36.446	39.652	49.088	1.00	37.75
ATOM	2550	O	ASN A 332	36.688	40.760	48.661	1.00	37.42
ATOM	2551	CB	ASN A 332	36.728	40.152	51.523	1.00	38.50
ATOM	2552	CG	ASN A 332	35.272	40.103	51.896	1.00	40.34
ATOM	2553	OD1	ASN A 332	34.493	39.377	51.274	1.00	43.85
ATOM	2554	ND2	ASN A 332	34.894	40.841	52.937	1.00	41.75
ATOM	2555	N	PRO A 333	35.628	38.839	48.451	1.00	37.92
ATOM	2556	CA	PRO A 333	35.076	39.209	47.146	1.00	37.97
ATOM	2557	C	PRO A 333	34.245	40.456	47.211	1.00	37.96
ATOM	2558	O	PRO A 333	34.086	41.143	46.204	1.00	37.81
ATOM	2559	CB	PRO A 333	34.182	38.024	46.770	1.00	37.85
ATOM	2560	CG	PRO A 333	34.327	37.037	47.801	1.00	38.26
ATOM	2561	CD	PRO A 333	35.161	37.530	48.908	1.00	37.98
ATOM	2562	N	GLN A 334	33.702	40.741	48.381	1.00	38.23
ATOM	2563	CA	GLN A 334	32.872	41.928	48.539	1.00	38.84
ATOM	2564	C	GLN A 334	33.713	43.177	48.442	1.00	37.04
ATOM	2565	O	GLN A 334	33.185	44.254	48.276	1.00	37.42
ATOM	2566	CB	GLN A 334	32.041	41.870	49.850	1.00	39.97
ATOM	2567	CG	GLN A 334	30.696	41.057	49.612	1.00	44.72
ATOM	2568	CD	GLN A 334	29.669	41.117	50.756	1.00	49.62
ATOM	2569	OE1	GLN A 334	29.666	42.067	51.566	1.00	53.65
ATOM	2570	NE2	GLN A 334	28.781	40.106	50.808	1.00	50.33
ATOM	2571	N	GLU A 335	35.026	43.036	48.489	1.00	35.15
ATOM	2572	CA	GLU A 335	35.872	44.193	48.385	1.00	34.55
ATOM	2573	C	GLU A 335	36.197	44.510	46.927	1.00	33.05
ATOM	2574	O	GLU A 335	36.760	45.564	46.640	1.00	32.79
ATOM	2575	CB	GLU A 335	37.144	44.028	49.247	1.00	35.19
ATOM	2576	CG	GLU A 335	36.927	44.326	50.727	1.00	37.92
ATOM	2577	CD	GLU A 335	38.195	44.232	51.550	1.00	43.46
ATOM	2578	OE1	GLU A 335	39.179	44.916	51.179	1.00	46.92
ATOM	2579	OE2	GLU A 335	38.211	43.499	52.589	1.00	49.17
ATOM	2580	N	VAL A 336	35.792	43.633	46.005	1.00	31.25
ATOM	2581	CA	VAL A 336	36.081	43.810	44.579	1.00	29.91
ATOM	2582	C	VAL A 336	35.505	45.076	43.957	1.00	29.08

C
C
O
C
C
C
O
O
N
C
C
O
C
N
C
C
O
C
C
C
N
C
C
O
N
C
C
C
O
C
C
C
N
C
C
C
O
C
C
C
O
C
C
O
N
N
C
C
C
O
C
C
O
C
O
N
C
C

ATOM	2583	O	VAL	A	336	36.177	45.816	43.246	1.00	29.08
ATOM	2584	CB	VAL	A	336	35.648	42.584	43.791	1.00	30.21
ATOM	2585	CG1	VAL	A	336	35.743	42.805	42.325	1.00	29.11
ATOM	2586	CG2	VAL	A	336	36.540	41.391	44.161	1.00	29.98
ATOM	2587	N	GLY	A	337	34.263	45.354	44.240	1.00	28.54
ATOM	2588	CA	GLY	A	337	33.644	46.546	43.722	1.00	27.74
ATOM	2589	C	GLY	A	337	34.285	47.848	44.130	1.00	27.24
ATOM	2590	O	GLY	A	337	34.599	48.687	43.281	1.00	26.81
ATOM	2591	N	PRO	A	338	34.365	48.094	45.428	1.00	27.19
ATOM	2592	CA	PRO	A	338	34.994	49.334	45.905	1.00	26.74
ATOM	2593	C	PRO	A	338	36.378	49.553	45.328	1.00	26.42
ATOM	2594	O	PRO	A	338	36.678	50.668	44.932	1.00	27.57
ATOM	2595	CB	PRO	A	338	35.008	49.173	47.426	1.00	25.92
ATOM	2596	CG	PRO	A	338	33.800	48.384	47.682	1.00	27.15
ATOM	2597	CD	PRO	A	338	33.750	47.326	46.531	1.00	27.09
ATOM	2598	N	LEU	A	339	37.199	48.533	45.225	1.00	25.95
ATOM	2599	CA	LEU	A	339	38.505	48.762	44.652	1.00	26.15
ATOM	2600	C	LEU	A	339	38.384	49.205	43.192	1.00	25.74
ATOM	2601	O	LEU	A	339	39.021	50.203	42.784	1.00	23.97
ATOM	2602	CB	LEU	A	339	39.375	47.521	44.763	1.00	27.19
ATOM	2603	CG	LEU	A	339	40.835	47.787	44.367	1.00	28.71
ATOM	2604	CD1	LEU	A	339	41.805	47.124	45.241	1.00	31.14
ATOM	2605	CD2	LEU	A	339	41.028	47.264	42.987	1.00	30.12
ATOM	2606	N	LEU	A	340	37.555	48.491	42.416	1.00	25.64
ATOM	2607	CA	LEU	A	340	37.364	48.841	40.991	1.00	25.90
ATOM	2608	C	LEU	A	340	36.860	50.261	40.871	1.00	26.04
ATOM	2609	O	LEU	A	340	37.324	51.044	40.034	1.00	25.93
ATOM	2610	CB	LEU	A	340	36.399	47.900	40.293	1.00	25.64
ATOM	2611	CG	LEU	A	340	36.996	46.521	40.072	1.00	27.50
ATOM	2612	CD1	LEU	A	340	35.943	45.559	39.555	1.00	26.88
ATOM	2613	CD2	LEU	A	340	38.221	46.587	39.138	1.00	26.63
ATOM	2614	N	ASN	A	341	35.914	50.610	41.720	1.00	26.45
ATOM	2615	CA	ASN	A	341	35.377	51.963	41.669	1.00	27.35
ATOM	2616	C	ASN	A	341	36.450	52.983	41.983	1.00	27.08
ATOM	2617	O	ASN	A	341	36.578	54.012	41.324	1.00	26.82
ATOM	2618	CB	ASN	A	341	34.183	52.117	42.619	1.00	27.50
ATOM	2619	CG	ASN	A	341	32.901	51.709	41.975	1.00	28.85
ATOM	2620	OD1	ASN	A	341	32.509	52.283	40.957	1.00	37.23
ATOM	2621	ND2	ASN	A	341	32.263	50.697	42.504	1.00	29.29
ATOM	2622	N	THR	A	342	37.234	52.665	42.994	1.00	26.89
ATOM	2623	CA	THR	A	342	38.321	53.509	43.355	1.00	27.22
ATOM	2624	C	THR	A	342	39.301	53.629	42.191	1.00	27.16

ATOM	2644	CD1	ILE	A	344	34.724	53.358	36.590	1.00	35.46	C
ATOM	2645	N	LYS	A	345	37.107	56.032	39.108	1.00	27.79	N
ATOM	2646	CA	LYS	A	345	36.518	57.364	39.218	1.00	28.48	C
ATOM	2647	C	LYS	A	345	37.489	58.485	39.219	1.00	27.72	C
ATOM	2648	O	LYS	A	345	38.428	58.555	40.000	1.00	27.37	O
ATOM	2649	CB	LYS	A	345	35.542	57.611	40.343	1.00	29.78	C
ATOM	2650	CG	LYS	A	345	35.537	56.758	41.491	1.00	34.49	C
ATOM	2651	CD	LYS	A	345	34.075	56.352	41.692	1.00	37.43	C
ATOM	2652	CE	LYS	A	345	33.508	57.036	42.904	1.00	40.47	C
ATOM	2653	NZ	LYS	A	345	34.223	56.522	44.138	1.00	46.19	N
ATOM	2654	N	GLY	A	346	37.218	59.386	38.297	1.00	26.82	N
ATOM	2655	CA	GLY	A	346	38.044	60.538	38.108	1.00	25.95	C
ATOM	2656	C	GLY	A	346	39.314	60.175	37.392	1.00	25.08	C
ATOM	2657	O	GLY	A	346	40.172	61.022	37.244	1.00	24.91	O
ATOM	2658	N	ARG	A	347	39.419	58.935	36.944	1.00	24.84	N
ATOM	2659	CA	ARG	A	347	40.647	58.436	36.347	1.00	25.30	C
ATOM	2660	C	ARG	A	347	40.431	57.848	34.937	1.00	26.05	C
ATOM	2661	O	ARG	A	347	41.138	58.196	34.007	1.00	24.59	O
ATOM	2662	CB	ARG	A	347	41.279	57.395	37.272	1.00	25.27	C
ATOM	2663	CG	ARG	A	347	41.716	57.934	38.696	1.00	24.21	C
ATOM	2664	CD	ARG	A	347	43.159	57.506	39.082	1.00	25.10	C
ATOM	2665	NE	ARG	A	347	43.175	56.084	39.017	1.00	24.49	N
ATOM	2666	CZ	ARG	A	347	44.102	55.293	38.566	1.00	19.51	C
ATOM	2667	NH1	ARG	A	347	45.299	55.692	38.195	1.00	21.86	N
ATOM	2668	NH2	ARG	A	347	43.802	54.011	38.579	1.00	18.94	N
ATOM	2669	N	TYR	A	348	39.438	56.991	34.794	1.00	27.91	N
ATOM	2670	CA	TYR	A	348	39.119	56.380	33.507	1.00	29.80	C
ATOM	2671	C	TYR	A	348	37.671	56.590	33.065	1.00	32.00	C
ATOM	2672	O	TYR	A	348	37.309	56.173	31.964	1.00	32.29	O
ATOM	2673	CB	TYR	A	348	39.379	54.877	33.543	1.00	28.38	C
ATOM	2674	CG	TYR	A	348	40.836	54.498	33.597	1.00	27.86	C
ATOM	2675	CD1	TYR	A	348	41.617	54.473	32.446	1.00	26.79	C
ATOM	2676	CD2	TYR	A	348	41.422	54.124	34.787	1.00	25.32	C
ATOM	2677	CE1	TYR	A	348	42.945	54.106	32.496	1.00	25.35	C
ATOM	2678	CE2	TYR	A	348	42.734	53.762	34.847	1.00	27.07	C
ATOM	2679	CZ	TYR	A	348	43.507	53.759	33.700	1.00	26.40	C
ATOM	2680	OH	TYR	A	348	44.827	53.378	33.789	1.00	22.93	O
ATOM	2681	N	ASN	A	349	36.824	57.193	33.894	1.00	34.78	N
ATOM	2682	CA	ASN	A	349	35.426	57.360	33.473	1.00	37.16	C
ATOM	2683	C	ASN	A	349	34.997	58.760	33.104	1.00	38.90	C
ATOM	2684	O	ASN	A	349	35.690	59.756	33.136	1.00	38.66	O
ATOM	2685	CB	ASN	A	349	34.474	56.837	34.493	1.00	36.69	C
ATOM	2686	CG	ASN	A	349	34.480	57.633	35.727	1.00	39.34	C
ATOM	2687	OD1	ASN	A	349	35.229	58.632	35.875	1.00	40.92	O
ATOM	2688	ND2	ASN	A	349	33.654	57.188	36.684	1.00	42.73	N
ATOM	2689	OXT	ASN	A	349	33.854	58.954	32.689	1.00	44.16	O
TER	2690		ASN	A	349						
ATOM	2691	N	LEU	S	795	45.870	35.442	31.163	1.00	49.68	N
ATOM	2692	CA	LEU	S	795	44.790	36.472	31.262	1.00	49.90	C
ATOM	2693	C	LEU	S	795	43.668	36.155	30.294	1.00	50.12	C
ATOM	2694	O	LEU	S	795	42.492	36.182	30.666	1.00	49.41	O
ATOM	2695	CB	LEU	S	795	45.331	37.873	31.004	1.00	50.09	C
ATOM	2696	CG	LEU	S	795	44.774	38.924	31.976	1.00	50.19	C
ATOM	2697	CD1	LEU	S	795	44.929	38.464	33.388	1.00	50.50	C
ATOM	2698	CD2	LEU	S	795	45.436	40.280	31.876	1.00	51.23	C
ATOM	2699	N	THR	S	796	44.041	35.979	29.033	1.00	50.69	N
ATOM	2700	CA	THR	S	796	43.178	35.421	27.997	1.00	51.67	C
ATOM	2701	C	THR	S	796	42.734	33.983	28.241	1.00	51.11	C
ATOM	2702	O	THR	S	796	41.884	33.464	27.535	1.00	51.41	O
ATOM	2703	CB	THR	S	796	43.888	35.547	26.626	1.00	51.86	C
ATOM	2704	OG1	THR	S	796	43.253	34.704	25.671	1.00	56.26	O

ATOM	2705	CG2	THR	S	796	45.272	34.981	26.662	1.00	52.69
ATOM	2706	N	SER	S	797	43.261	33.330	29.248	1.00	51.51
ATOM	2707	CA	SER	S	797	42.838	31.965	29.497	1.00	52.74
ATOM	2708	C	SER	S	797	41.471	31.865	30.165	1.00	53.39
ATOM	2709	O	SER	S	797	41.070	32.733	30.934	1.00	52.91
ATOM	2710	CB	SER	S	797	43.850	31.247	30.351	1.00	52.86
ATOM	2711	OG	SER	S	797	43.687	31.623	31.697	1.00	55.21
ATOM	2712	N	TYR	S	798	40.783	30.761	29.904	1.00	54.79
ATOM	2713	CA	TYR	S	798	39.424	30.584	30.372	1.00	56.02
ATOM	2714	C	TYR	S	798	39.168	29.271	31.042	1.00	55.81
ATOM	2715	O	TYR	S	798	39.880	28.305	30.827	1.00	56.61
ATOM	2716	CB	TYR	S	798	38.437	30.747	29.214	1.00	56.97
ATOM	2717	CG	TYR	S	798	38.599	29.821	28.015	1.00	60.68
ATOM	2718	CD1	TYR	S	798	39.595	30.035	27.059	1.00	62.78
ATOM	2719	CD2	TYR	S	798	37.689	28.783	27.793	1.00	64.68
ATOM	2720	CE1	TYR	S	798	39.708	29.207	25.943	1.00	65.08
ATOM	2721	CE2	TYR	S	798	37.798	27.944	26.669	1.00	66.15
ATOM	2722	CZ	TYR	S	798	38.808	28.155	25.755	1.00	65.29
ATOM	2723	OH	TYR	S	798	38.902	27.335	24.652	1.00	64.08
ATOM	2724	N	ASP	S	799	38.117	29.240	31.843	1.00	55.66
ATOM	2725	CA	ASP	S	799	37.727	28.032	32.546	1.00	55.90
ATOM	2726	C	ASP	S	799	36.636	27.385	31.739	1.00	54.64
ATOM	2727	O	ASP	S	799	36.430	27.785	30.604	1.00	55.18
ATOM	2728	CB	ASP	S	799	37.212	28.367	33.948	1.00	56.52
ATOM	2729	CG	ASP	S	799	37.366	27.225	34.906	1.00	57.36
ATOM	2730	OD1	ASP	S	799	36.749	26.147	34.720	1.00	57.64
ATOM	2731	OD2	ASP	S	799	38.108	27.331	35.881	1.00	62.28
ATOM	2732	N	CYS	S	800	35.915	26.428	32.326	1.00	53.45
ATOM	2733	CA	CYS	S	800	34.887	25.693	31.594	1.00	52.63
ATOM	2734	C	CYS	S	800	33.564	25.644	32.359	1.00	51.79
ATOM	2735	O	CYS	S	800	32.871	24.632	32.357	1.00	51.30
ATOM	2736	CB	CYS	S	800	35.372	24.282	31.327	1.00	52.47
ATOM	2737	SG	CYS	S	800	35.703	23.408	32.872	1.00	52.72
ATOM	2738	N	GLU	S	801	33.187	26.742	32.996	1.00	50.91
ATOM	2739	CA	GLU	S	801	31.939	26.732	33.762	1.00	50.65
ATOM	2740	C	GLU	S	801	30.657	26.939	32.909	1.00	50.18
ATOM	2741	O	GLU	S	801	30.617	27.687	31.928	1.00	48.39
ATOM	2742	CB	GLU	S	801	32.008	27.713	34.929	1.00	50.49
ATOM	2743	CG	GLU	S	801	33.199	27.450	35.853	1.00	51.78
ATOM	2744	CD	GLU	S	801	33.234	28.359	37.092	1.00	51.24
ATOM	2745	OE1	GLU	S	801	32.589	28.038	38.102	1.00	49.25
ATOM	2746	OE2	GLU	S	801	33.919	29.395	37.063	1.00	51.89
ATOM	2747	N	VAL	S	802	29.624	26.213	33.320	1.00	50.54
ATOM	2748	CA	VAL	S	802	28.338	26.161	32.650	1.00	50.71
ATOM	2749	C	VAL	S	802	27.213	26.094	33.694	1.00	51.25
ATOM	2750	O	VAL	S	802	27.464	25.960	34.885	1.00	50.69
ATOM	2751	CB	VAL	S	802	28.286	24.902	31.746	1.00	50.61
ATOM	2752	CG1	VAL	S	802	29.420	24.932	30.727	1.00	49.62
ATOM	2753	CG2	VAL	S	802	28.376	23.601	32.585	1.00	49.78
ATOM	2754	N	ASN	S	803	25.967	26.165	33.239	1.00	52.55
ATOM	2755	CA	ASN	S	803	24.816	26.099	34.140	1.00	53.28
ATOM	2756	C	ASN	S	803	24.516	24.663	34.561	1.00	54.95
ATOM	2757	O	ASN	S	803	23.440	24.150	34.298	1.00	54.22
ATOM	2758	CB	ASN	S	803	23.560	26.724	33.511	1.00	53.04
ATOM	2759	CG	ASN	S	803	23.546	28.245	33.569	1.00	51.60
ATOM	2760	OD1	ASN	S	803	24.576	28.895	33.755	1.00	51.13
ATOM	2761	ND2	ASN	S	803	22.367	28.817	33.406	1.00	46.28
ATOM	2762	N	ALA	S	804	25.498	24.033	35.201	1.00	57.29
ATOM	2763	CA	ALA	S	804	25.373	22.697	35.784	1.00	59.38
ATOM	2764	C	ALA	S	804	26.596	22.490	36.676	1.00	61.01
ATOM	2765	O	ALA	S	804	27.667	23.026	36.407	1.00	61.45

C
N
C
C
O
C
O
N
C
C
C
C
C
C
O
N
C
C
C
O
C
S
N
C
C
O
C
C
C
O
O
N
C
C
C
C
O
N
N
C
C
O

ATOM	2766	CB	ALA	S	804	25.315	21.603	34.716	1.00	59.47
ATOM	2767	N	PRO	S	805	26.438	21.707	37.732	1.00	62.94
ATOM	2768	CA	PRO	S	805	27.547	21.398	38.652	1.00	63.79
ATOM	2769	C	PRO	S	805	28.616	20.498	38.003	1.00	64.48
ATOM	2770	O	PRO	S	805	28.368	19.875	36.970	1.00	65.26
ATOM	2771	CB	PRO	S	805	26.851	20.654	39.807	1.00	63.79
ATOM	2772	CG	PRO	S	805	25.582	20.066	39.187	1.00	63.48
ATOM	2773	CD	PRO	S	805	25.175	21.035	38.107	1.00	63.26
ATOM	2774	N	ILE	S	806	29.779	20.405	38.630	1.00	65.18
ATOM	2775	CA	ILE	S	806	30.903	19.626	38.100	1.00	65.66
ATOM	2776	C	ILE	S	806	30.853	18.145	38.489	1.00	65.84
ATOM	2777	O	ILE	S	806	30.287	17.780	39.522	1.00	66.22
ATOM	2778	CB	ILE	S	806	32.239	20.255	38.582	1.00	65.91
ATOM	2779	CG1	ILE	S	806	32.435	20.046	40.089	1.00	65.62
ATOM	2780	CG2	ILE	S	806	32.267	21.765	38.255	1.00	66.46
ATOM	2781	CD1	ILE	S	806	33.719	20.639	40.611	1.00	65.48
ATOM	2782	N	LEU	S	813	29.912	8.313	36.265	1.00	60.30
ATOM	2783	CA	LEU	S	813	30.605	8.781	35.059	1.00	60.57
ATOM	2784	C	LEU	S	813	29.707	9.617	34.157	1.00	60.31
ATOM	2785	O	LEU	S	813	28.484	9.496	34.194	1.00	60.27
ATOM	2786	CB	LEU	S	813	31.167	7.599	34.266	1.00	60.46
ATOM	2787	CG	LEU	S	813	32.067	6.674	35.092	1.00	60.30
ATOM	2788	CD1	LEU	S	813	32.450	5.418	34.281	1.00	61.04
ATOM	2789	CD2	LEU	S	813	33.301	7.421	35.602	1.00	59.11
ATOM	2790	N	GLN	S	814	30.344	10.465	33.355	1.00	60.31
ATOM	2791	CA	GLN	S	814	29.648	11.353	32.440	1.00	60.05
ATOM	2792	C	GLN	S	814	30.557	11.803	31.295	1.00	59.62
ATOM	2793	O	GLN	S	814	31.765	11.584	31.322	1.00	58.97
ATOM	2794	CB	GLN	S	814	29.176	12.583	33.194	1.00	60.41
ATOM	2795	CG	GLN	S	814	30.302	13.488	33.634	1.00	62.13
ATOM	2796	CD	GLN	S	814	29.826	14.603	34.559	1.00	64.43
ATOM	2797	OE1	GLN	S	814	29.260	14.327	35.616	1.00	65.48
ATOM	2798	NE2	GLN	S	814	30.058	15.859	34.167	1.00	65.06
ATOM	2799	N	GLY	S	815	29.960	12.456	30.302	1.00	59.55
ATOM	2800	CA	GLY	S	815	30.675	12.927	29.136	1.00	59.21
ATOM	2801	C	GLY	S	815	31.493	11.823	28.493	1.00	59.26
ATOM	2802	O	GLY	S	815	31.049	10.678	28.399	1.00	58.48
ATOM	2803	N	GLU	S	816	32.693	12.183	28.049	1.00	59.73
ATOM	2804	CA	GLU	S	816	33.604	11.254	27.392	1.00	60.76
ATOM	2805	C	GLU	S	816	33.768	9.941	28.168	1.00	61.61
ATOM	2806	O	GLU	S	816	33.861	8.866	27.567	1.00	61.64
ATOM	2807	CB	GLU	S	816	34.978	11.913	27.190	1.00	60.71
ATOM	2808	CG	GLU	S	816	35.802	11.304	26.064	1.00	61.12
ATOM	2809	CD	GLU	S	816	37.208	11.872	25.979	1.00	61.35
ATOM	2810	OE1	GLU	S	816	38.066	11.370	26.730	1.00	61.67
ATOM	2811	OE2	GLU	S	816	37.467	12.801	25.168	1.00	60.68
ATOM	2812	N	GLU	S	817	33.791	10.030	29.495	1.00	62.56
ATOM	2813	CA	GLU	S	817	33.962	8.851	30.332	1.00	63.66
ATOM	2814	C	GLU	S	817	32.748	7.929	30.260	1.00	63.92
ATOM	2815	O	GLU	S	817	32.887	6.709	30.187	1.00	63.68
ATOM	2816	CB	GLU	S	817	34.217	9.257	31.777	1.00	63.99
ATOM	2817	CG	GLU	S	817	35.518	10.012	31.974	1.00	65.72
ATOM	2818	CD	GLU	S	817	35.371	11.514	31.786	1.00	69.17
ATOM	2819	OE1	GLU	S	817	34.253	11.990	31.452	1.00	70.86
ATOM	2820	OE2	GLU	S	817	36.384	12.228	31.984	1.00	71.76
ATOM	2821	N	LEU	S	818	31.558	8.517	30.296	1.00	64.38
ATOM	2822	CA	LEU	S	818	30.344	7.733	30.191	1.00	65.00
ATOM	2823	C	LEU	S	818	30.367	6.935	28.890	1.00	65.31
ATOM	2824	O	LEU	S	818	30.166	5.723	28.901	1.00	65.41
ATOM	2825	CB	LEU	S	818	29.092	8.624	30.232	1.00	64.90
ATOM	2826	CG	LEU	S	818	27.770	7.844	30.163	1.00	64.94

C
N
C
C
O
C
C
N
C
C
O
C
C
N
C
C
O
C
C
C
C
N
C
C
O
C
C
O
N
N
C
C
O
N
C
C
C
O
C
C
O
O
N
C
C
C
O
O
N
C
C
C
C

ATOM	2827	CD1	LEU	S	818	27.702	6.871	31.316	1.00	66.53	
ATOM	2828	CD2	LEU	S	818	26.551	8.718	30.205	1.00	63.99	C
ATOM	2829	N	LEU	S	819	30.647	7.627	27.787	1.00	65.71	C
ATOM	2830	CA	LEU	S	819	30.608	7.037	26.445	1.00	66.10	C
ATOM	2831	C	LEU	S	819	31.569	5.851	26.294	1.00	66.37	C
ATOM	2832	O	LEU	S	819	31.174	4.773	25.852	1.00	65.86	O
ATOM	2833	CB	LEU	S	819	30.901	8.122	25.392	1.00	66.04	C
ATOM	2834	CG	LEU	S	819	30.860	7.737	23.916	1.00	66.00	C
ATOM	2835	CD1	LEU	S	819	29.537	7.149	23.501	1.00	66.14	C
ATOM	2836	CD2	LEU	S	819	31.162	8.967	23.083	1.00	67.31	C
ATOM	2837	N	ARG	S	820	32.820	6.053	26.684	1.00	66.68	C
ATOM	2838	CA	ARG	S	820	33.824	5.013	26.561	1.00	67.34	N
ATOM	2839	C	ARG	S	820	33.492	3.801	27.432	1.00	67.41	C
ATOM	2840	O	ARG	S	820	33.573	2.643	26.969	1.00	67.19	C
ATOM	2841	CB	ARG	S	820	35.196	5.574	26.917	1.00	67.66	C
ATOM	2842	CG	ARG	S	820	35.628	6.605	25.907	1.00	69.12	C
ATOM	2843	CD	ARG	S	820	37.048	7.061	26.030	1.00	71.00	C
ATOM	2844	NE	ARG	S	820	37.309	8.138	25.079	1.00	73.46	C
ATOM	2845	CZ	ARG	S	820	38.442	8.839	25.014	1.00	75.31	N
ATOM	2846	NH1	ARG	S	820	39.451	8.581	25.851	1.00	75.90	C
ATOM	2847	NH2	ARG	S	820	38.566	9.806	24.104	1.00	75.55	N
ATOM	2848	N	ALA	S	821	33.118	4.067	28.684	1.00	67.11	N
ATOM	2849	CA	ALA	S	821	32.712	2.999	29.580	1.00	66.99	N
ATOM	2850	C	ALA	S	821	31.609	2.171	28.898	1.00	66.76	C
ATOM	2851	O	ALA	S	821	31.660	0.945	28.889	1.00	66.63	C
ATOM	2852	CB	ALA	S	821	32.234	3.562	30.901	1.00	66.88	O
ATOM	2853	N	LEU	S	822	30.625	2.851	28.313	1.00	66.58	C
ATOM	2854	CA	LEU	S	822	29.538	2.173	27.607	1.00	66.36	N
ATOM	2855	C	LEU	S	822	30.027	1.439	26.359	1.00	66.21	C
ATOM	2856	O	LEU	S	822	29.429	0.444	25.956	1.00	66.11	C
ATOM	2857	CB	LEU	S	822	28.451	3.172	27.223	1.00	66.06	O
ATOM	2858	CG	LEU	S	822	27.710	3.789	28.406	1.00	65.45	C
ATOM	2859	CD1	LEU	S	822	26.559	4.613	27.885	1.00	65.37	C
ATOM	2860	CD2	LEU	S	822	27.213	2.732	29.396	1.00	65.08	C
TER	2861		LEU	S	822						
HETATM	2862	ZN	ZN	A1350		23.324	27.578	28.817	1.00	32.29	ZN
HETATM	2863	C1	OGA	A1351		22.262	25.308	27.891	1.00	34.79	C
HETATM	2864	C2	OGA	A1351		21.209	25.940	28.487	1.00	32.90	C
HETATM	2865	C4	OGA	A1351		18.882	25.730	29.253	1.00	32.24	C
HETATM	2866	C5	OGA	A1351		17.543	25.251	28.735	1.00	32.20	C
HETATM	2867	O1	OGA	A1351		22.091	24.207	27.399	1.00	33.47	O
HETATM	2868	O2	OGA	A1351		23.404	25.817	27.813	1.00	33.83	O
HETATM	2869	O2'	OGA	A1351		21.299	27.077	29.004	1.00	31.02	O
HETATM	2870	O3	OGA	A1351		17.430	24.476	27.795	1.00	33.67	O
HETATM	2871	N1	OGA	A1351		20.067	25.271	28.520	1.00	29.83	N
HETATM	2872	O4	OGA	A1351		16.551	25.641	29.307	1.00	32.21	O
HETATM	2873	S	SO4	A1352		0.290	25.194	43.827	1.00	90.02	S
HETATM	2874	O1	SO4	A1352		1.120	26.025	44.689	1.00	89.95	O
HETATM	2875	O2	SO4	A1352		1.151	24.261	43.106	1.00	88.91	O
HETATM	2876	O3	SO4	A1352		-0.627	24.447	44.672	1.00	90.38	O
HETATM	2877	O4	SO4	A1352		-0.468	26.028	42.891	1.00	89.30	O
HETATM	2878	S	SO4	A1353		1.893	28.515	29.870	1.00	98.62	S
HETATM	2879	O1	SO4	A1353		3.138	29.102	30.350	1.00	97.69	O
HETATM	2880	O2	SO4	A1353		2.145	27.399	28.947	1.00	97.91	O
HETATM	2881	O3	SO4	A1353		1.205	28.059	31.078	1.00	99.26	O
HETATM	2882	O4	SO4	A1353		1.078	29.515	29.171	1.00	98.46	O
HETATM	2883	O	HOH	H	1	38.820	33.858	31.965	1.00	46.43	O
HETATM	2884	O	HOH	H	2	33.795	30.509	39.255	1.00	71.93	O
HETATM	2885	O	HOH	H	3	34.891	30.536	35.372	1.00	48.26	O
HETATM	2886	O	HOH	H	4	35.615	13.844	24.220	1.00	48.67	O
HETATM	2887	O	HOH	Z	1	11.592	21.463	13.878	1.00	49.06	O

HETATM	2949	O	HOH	Z	63	26.759	31.212	38.332	1.00	39.21	O
HETATM	2950	O	HOH	Z	64	29.563	36.988	35.538	1.00	36.58	O
HETATM	2951	O	HOH	Z	65	30.608	30.268	36.508	1.00	39.97	O
HETATM	2952	O	HOH	Z	66	19.823	31.620	33.966	1.00	44.64	O
HETATM	2953	O	HOH	Z	67	19.517	36.182	33.531	1.00	39.38	O
HETATM	2954	O	HOH	Z	68	18.255	38.678	36.813	1.00	57.99	O
HETATM	2955	O	HOH	Z	69	21.026	42.838	40.240	1.00	56.11	O
HETATM	2956	O	HOH	Z	70	17.610	37.832	32.310	1.00	47.08	O
HETATM	2957	O	HOH	Z	71	15.109	37.973	32.827	1.00	45.82	O
HETATM	2958	O	HOH	Z	72	16.418	47.375	39.219	1.00	67.81	O
HETATM	2959	O	HOH	Z	73	15.605	44.895	35.040	1.00	43.78	O
HETATM	2960	O	HOH	Z	74	15.286	55.715	27.660	1.00	67.46	O
HETATM	2961	O	HOH	Z	75	15.476	50.403	22.363	1.00	44.42	O
HETATM	2962	O	HOH	Z	76	16.645	46.682	22.198	1.00	39.25	O
HETATM	2963	O	HOH	Z	77	26.589	40.795	15.622	1.00	68.21	O
HETATM	2964	O	HOH	Z	78	10.674	19.290	14.058	1.00	55.87	O
HETATM	2965	O	HOH	Z	79	27.773	27.556	24.476	1.00	37.12	O
HETATM	2966	O	HOH	Z	80	30.214	36.062	27.011	1.00	40.11	O
HETATM	2967	O	HOH	Z	81	32.661	34.179	19.369	1.00	54.04	O
HETATM	2968	O	HOH	Z	82	27.339	30.221	25.948	1.00	38.07	O
HETATM	2969	O	HOH	Z	83	16.261	36.239	29.815	1.00	43.65	O
HETATM	2970	O	HOH	Z	84	6.400	23.973	25.697	1.00	45.37	O
HETATM	2971	O	HOH	Z	85	4.061	20.777	20.758	1.00	74.60	O
HETATM	2972	O	HOH	Z	86	42.841	37.277	18.598	1.00	67.00	O
HETATM	2973	O	HOH	Z	87	42.456	44.259	19.547	1.00	64.85	O
HETATM	2974	O	HOH	Z	88	33.750	36.936	20.231	1.00	46.70	O
HETATM	2975	O	HOH	Z	89	33.208	39.305	33.829	1.00	37.00	O
HETATM	2976	O	HOH	Z	90	32.098	40.251	44.432	1.00	46.81	O
HETATM	2977	O	HOH	Z	91	32.904	43.762	45.581	1.00	58.90	O
HETATM	2978	O	HOH	Z	92	39.242	57.294	42.204	1.00	36.08	O
HETATM	2979	O	HOH	Z	93	46.926	53.378	32.000	1.00	42.91	O
HETATM	2980	O	HOH	Z	94	37.922	55.476	29.664	1.00	46.42	O
HETATM	2981	O	HOH	Z	95	38.489	59.800	33.014	1.00	55.47	O
CONNECT	1478	2862									
CONNECT	1498	2862									
CONNECT	2167	2862									
CONNECT	2862	2869	2868	1478	2167	1498					
CONNECT	2863	2864	2867	2868							
CONNECT	2864	2863	2869	2871							
CONNECT	2865	2866	2871								
CONNECT	2866	2865	2870	2872							
CONNECT	2867	2863									
CONNECT	2868	2863	2862								

Structure 4

Below are the coordinates for structure 4 (the 2.85 Å structure of FIH:Fe(II):2OG):

```

HEADER    TRANSCRIPTION ACTIVATOR/INHIBITOR      12-AUG-02   1H2N
TITLE     FACTOR INHIBITING HIF-1 ALPHA IN COMPLEX WITH HIF-1 ALPHA
TITLE     2 FRAGMENT PEPTIDE
COMPND    MOL_ID: 1;
COMPND    2 MOLECULE: FACTOR INHIBITING HIF1;
COMPND    3 SYNONYM: FIH1;
COMPND    4 CHAIN: A;
COMPND    5 ENGINEERED: YES
SOURCE    MOL_ID: 1;
SOURCE    2 ORGANISM_SCIENTIFIC: HOMO SAPIENS;
SOURCE    3 ORGANISM_COMMON: HUMAN;
SOURCE    4 EXPRESSION_SYSTEM: ESCHERICHIA COLI;
SOURCE    5 EXPRESSION_SYSTEM_STRAIN: BL21(DE3);
SOURCE    6 EXPRESSION_SYSTEM_PLASMID: PET28A(+)
KEYWDS    FIH, HIF, DSBH, OXYGENASE, TRANSCRIPTION, HYPOXIA,
KEYWDS    2 2-OXOGLUTARATE, ASPARAGINYL HYDROXYLASE, HYDROXYLASE
EXPDTA    X-RAY DIFFRACTION
AUTHOR    J.M.ELKINS,K.S.HEWITSON,L.A.MCNEILL,I.SCHLEMMINGER,
AUTHOR    2 J.F.SEIBEL,C.J.SCHOFIELD
REVDAT    1 04-SEP-02 1H2N 0
JRNL      AUTH J.M.ELKINS,K.S.HEWITSON,L.A.MCNEILL,
JRNL      AUTH 2 I.SCHLEMMINGER,J.F.SEIBEL,C.J.SCHOFIELD
JRNL      TITL 2 FIH:HIF-FRAGMENT COMPLEXES
JRNL      REF  TO BE PUBLISHED
JRNL      REFN
REMARK    2
REMARK    2 RESOLUTION. 2.84 ANGSTROMS.
REMARK    3
REMARK    3 REFINEMENT.
REMARK    3 PROGRAM : REFMAC 5.0
REMARK    3 AUTHORS : MURSHUDOV,VAGIN,DODSON
REMARK    3
REMARK    3 REFINEMENT TARGET : MAXIMUM LIKELIHOOD
REMARK    3
REMARK    3 DATA USED IN REFINEMENT.
REMARK    3 RESOLUTION RANGE HIGH (ANGSTROMS) : 2.84
REMARK    3 RESOLUTION RANGE LOW (ANGSTROMS) : 30.00
REMARK    3 DATA CUTOFF (SIGMA(F)) : NONE
REMARK    3 COMPLETENESS FOR RANGE (%) : 99.51
REMARK    3 NUMBER OF REFLECTIONS : 12577
REMARK    3
REMARK    3 FIT TO DATA USED IN REFINEMENT.
REMARK    3 CROSS-VALIDATION METHOD : THROUGHOUT
REMARK    3 FREE R VALUE TEST SET SELECTION : RANDOM
REMARK    3 R VALUE (WORKING + TEST SET) : 0.23287
REMARK    3 R VALUE (WORKING SET) : 0.23094
REMARK    3 FREE R VALUE : 0.25695
REMARK    3 FREE R VALUE TEST SET SIZE (%) : 7.7
REMARK    3 FREE R VALUE TEST SET COUNT : 1046
REMARK    3
REMARK    3 FIT IN THE HIGHEST RESOLUTION BIN.
REMARK    3 TOTAL NUMBER OF BINS USED : 20
REMARK    3 BIN RESOLUTION RANGE HIGH : 2.840
REMARK    3 BIN RESOLUTION RANGE LOW : 2.913
REMARK    3 REFLECTION IN BIN (WORKING SET) : 828

```

REMARK 3 BIN R VALUE (WORKING SET) : 0.286
 REMARK 3 BIN FREE R VALUE SET COUNT : 81
 REMARK 3 BIN FREE R VALUE : 0.315
 REMARK 3
 REMARK 3 NUMBER OF NON-HYDROGEN ATOMS USED IN REFINEMENT.
 REMARK 3 PROTEIN ATOMS : 2689
 REMARK 3 NUCLEIC ACID ATOMS : 0
 REMARK 3 HETEROGEN ATOMS : 26
 REMARK 3 SOLVENT ATOMS : 3
 REMARK 3
 REMARK 3 B VALUES.
 REMARK 3 FROM WILSON PLOT (A**2) : NULL
 REMARK 3 MEAN B VALUE (OVERALL, A**2) : 35.345
 REMARK 3 OVERALL ANISOTROPIC B VALUE.
 REMARK 3 B11 (A**2) : -1.02
 REMARK 3 B22 (A**2) : -1.02
 REMARK 3 B33 (A**2) : 2.03
 REMARK 3 B12 (A**2) : 0.00
 REMARK 3 B13 (A**2) : 0.00
 REMARK 3 B23 (A**2) : 0.00
 REMARK 3
 REMARK 3 ESTIMATED OVERALL COORDINATE ERROR.
 REMARK 3 ESU BASED ON R VALUE (A): 0.852
 REMARK 3 ESU BASED ON FREE R VALUE (A): 0.349
 REMARK 3 ESU BASED ON MAXIMUM LIKELIHOOD (A): 0.398
 REMARK 3 ESU FOR B VALUES BASED ON MAXIMUM LIKELIHOOD (A**2): 19.679
 REMARK 3
 REMARK 3 CORRELATION COEFFICIENTS.
 REMARK 3 CORRELATION COEFFICIENT FO-FC : 0.913
 REMARK 3 CORRELATION COEFFICIENT FO-FC FREE : 0.901
 REMARK 3
 REMARK 3 RMS DEVIATIONS FROM IDEAL VALUES COUNT RMS WEIGHT
 REMARK 3 BOND LENGTHS REFINED ATOMS (A): 2791 ; 0.015 ; 0.021
 REMARK 3 BOND LENGTHS OTHERS (A): 2388 ; 0.001 ; 0.020
 REMARK 3 BOND ANGLES REFINED ATOMS (DEGREES): 3799 ; 1.628 ; 1.945
 REMARK 3 BOND ANGLES OTHERS (DEGREES): 5576 ; 0.823 ; 3.000
 REMARK 3 TORSION ANGLES, PERIOD 1 (DEGREES): 330 ; 4.268 ; 3.000
 REMARK 3 TORSION ANGLES, PERIOD 3 (DEGREES): 479 ; 18.082 ; 15.000
 REMARK 3 CHIRAL-CENTER RESTRAINTS (A**3): 384 ; 0.095 ; 0.200
 REMARK 3 GENERAL PLANES REFINED ATOMS (A): 3137 ; 0.005 ; 0.020
 REMARK 3 GENERAL PLANES OTHERS (A): 575 ; 0.002 ; 0.020
 REMARK 3 NON-BONDED CONTACTS REFINED ATOMS (A): 717 ; 0.255 ; 0.300
 REMARK 3 NON-BONDED CONTACTS OTHERS (A): 2425 ; 0.224 ; 0.300
 REMARK 3 H-BOND (X...Y) REFINED ATOMS (A): 165 ; 0.139 ; 0.500
 REMARK 3 H-BOND (X...Y) OTHERS (A): 1 ; 0.102 ; 0.500
 REMARK 3 POTENTIAL METAL-ION REFINED ATOMS (A): 3 ; 0.112 ; 0.500
 REMARK 3 SYMMETRY VDW REFINED ATOMS (A): 14 ; 0.256 ; 0.300
 REMARK 3 SYMMETRY VDW OTHERS (A): 62 ; 0.273 ; 0.300
 REMARK 3 SYMMETRY H-BOND REFINED ATOMS (A): 4 ; 0.214 ; 0.500
 REMARK 3 SYMMETRY H-BOND OTHERS (A): 1 ; 0.061 ; 0.500
 REMARK 3
 REMARK 3 ISOTROPIC THERMAL FACTOR RESTRAINTS. COUNT RMS WEIGHT
 REMARK 3 MAIN-CHAIN BOND REFINED ATOMS (A**2): 1659 ; 0.312 ; 1.500
 REMARK 3 MAIN-CHAIN ANGLE REFINED ATOMS (A**2): 2675 ; 0.598 ; 2.000
 REMARK 3 SIDE-CHAIN BOND REFINED ATOMS (A**2): 1132 ; 1.058 ; 3.000
 REMARK 3 SIDE-CHAIN ANGLE REFINED ATOMS (A**2): 1124 ; 1.795 ; 4.500
 REMARK 3
 REMARK 3 NCS RESTRAINTS STATISTICS
 REMARK 3 NUMBER OF NCS GROUPS : NULL
 REMARK 3
 REMARK 3 TLS DETAILS

REMARK 3 NUMBER OF TLS GROUPS : 1
REMARK 3
REMARK 3 TLS GROUP : 1
REMARK 3 NUMBER OF COMPONENTS GROUP : 1
REMARK 3 COMPONENTS C SSSEQI TO C SSSEQI
REMARK 3 RESIDUE RANGE : A 15 A 452
REMARK 3 ORIGIN FOR THE GROUP (A): 21.4490 27.4200 27.7870
REMARK 3 T TENSOR
REMARK 3 T11: 0.2230 T22: 0.0562
REMARK 3 T33: 0.0967 T12: 0.0111
REMARK 3 T13: -0.0923 T23: 0.0525
REMARK 3 L TENSOR
REMARK 3 L11: 1.6842 L22: 4.4489
REMARK 3 L33: 2.0658 L12: 1.5597
REMARK 3 L13: 1.1572 L23: 2.3523
REMARK 3 S TENSOR
REMARK 3 S11: 0.1098 S12: -0.2106 S13: -0.0766
REMARK 3 S21: 0.3449 S22: -0.0455 S23: 0.2455
REMARK 3 S31: 0.3515 S32: -0.1199 S33: -0.0643
REMARK 3
REMARK 3 BULK SOLVENT MODELLING.
REMARK 3 METHOD USED : BABINET MODEL WITH MASK
REMARK 3 PARAMETERS FOR MASK CALCULATION
REMARK 3 VDW PROBE RADIUS : 1.40
REMARK 3 ION PROBE RADIUS : 0.80
REMARK 3 SHRINKAGE RADIUS : 0.80
REMARK 3
REMARK 3 OTHER REFINEMENT REMARKS: SEE REMARK 400
REMARK 4
REMARK 4 1H2N COMPLIES WITH FORMAT V. 2.3, 09-JULY-1998
REMARK 100
REMARK 100 THIS ENTRY HAS BEEN PROCESSED BY EBI ON 12-AUG-2002.
REMARK 100 THE EBI ID CODE IS EBI-11174.
REMARK 200
REMARK 200 EXPERIMENTAL DETAILS
REMARK 200 EXPERIMENT TYPE : X-RAY DIFFRACTION
REMARK 200 DATE OF DATA COLLECTION : 15-MAY-2002
REMARK 200 TEMPERATURE (KELVIN) : 100
REMARK 200 PH : 7.5
REMARK 200 NUMBER OF CRYSTALS USED : 1
REMARK 200
REMARK 200 SYNCHROTRON (Y/N) : Y
REMARK 200 RADIATION SOURCE : SRS BEAMLINE PX9.5
REMARK 200 BEAMLINE : PX9.5
REMARK 200 X-RAY GENERATOR MODEL : NULL
REMARK 200 MONOCHROMATIC OR LAUE (M/L) : M
REMARK 200 WAVELENGTH OR RANGE (A) : 0.92
REMARK 200 MONOCHROMATOR : NULL
REMARK 200 OPTICS : NULL
REMARK 200
REMARK 200 DETECTOR TYPE : MARCCD
REMARK 200 DETECTOR MANUFACTURER : MARRESEARCH
REMARK 200 INTENSITY-INTEGRATION SOFTWARE : MOSFLM
REMARK 200 DATA SCALING SOFTWARE : SCALA
REMARK 200
REMARK 200 NUMBER OF UNIQUE REFLECTIONS : 13703
REMARK 200 RESOLUTION RANGE HIGH (A) : 2.84
REMARK 200 RESOLUTION RANGE LOW (A) : 34.1
REMARK 200 REJECTION CRITERIA (SIGMA(I)) : NONE
REMARK 200
REMARK 200 OVERALL.

REMARK 200 COMPLETENESS FOR RANGE (%) : 99.2
 REMARK 200 DATA REDUNDANCY : 6.7
 REMARK 200 R MERGE (I) : 0.067
 REMARK 200 R SYM (I) : NULL
 REMARK 200 <I/SIGMA(I)> FOR THE DATA SET : 9.4
 REMARK 200
 REMARK 200 IN THE HIGHEST RESOLUTION SHELL.
 REMARK 200 HIGHEST RESOLUTION SHELL, RANGE HIGH (A) : 2.84
 REMARK 200 HIGHEST RESOLUTION SHELL, RANGE LOW (A) : 2.99
 REMARK 200 COMPLETENESS FOR SHELL (%) : 94.9
 REMARK 200 DATA REDUNDANCY IN SHELL : 5.2
 REMARK 200 R MERGE FOR SHELL (I) : 0.309
 REMARK 200 R SYM FOR SHELL (I) : NULL
 REMARK 200 <I/SIGMA(I)> FOR SHELL : 2.4
 REMARK 200
 REMARK 200 DIFFRACTION PROTOCOL: SINGLE WAVELENGTH
 REMARK 200 METHOD USED TO DETERMINE THE STRUCTURE: MOLECULAR REPLACEMENT
 REMARK 200 SOFTWARE USED: NULL
 REMARK 200 STARTING MODEL: NULL
 REMARK 200
 REMARK 200 REMARK: SEE REMARK 400
 REMARK 280
 REMARK 280 CRYSTAL
 REMARK 280 SOLVENT CONTENT, VS (%): 63
 REMARK 280 MATTHEWS COEFFICIENT, VM (ANGSTROMS**3/DA): 3.4
 REMARK 280
 REMARK 280 CRYSTALLIZATION CONDITIONS: 1.2M AMMONIUM SULPHATE,
 REMARK 280 4% PEG400, 0.1M HEPES PH7.5 ARGON ATMOSPHERE,
 REMARK 280 11MG/ML PROTEIN WITH 1MM FE(II), 2.5MM AKG AND 2.5MM
 REMARK 280 PEPTIDE (SEE REMARK 400)
 REMARK 290
 REMARK 290 CRYSTALLOGRAPHIC SYMMETRY
 REMARK 290 SYMMETRY OPERATORS FOR SPACE GROUP: P 41 21 2
 REMARK 290

SYNOP	SYMMETRY
NNNMMM	OPERATOR
1555	X, Y, Z
2555	-X, -Y, 1/2+Z
3555	1/2-Y, 1/2+X, 1/4+Z
4555	1/2+Y, 1/2-X, 3/4+Z
5555	1/2-X, 1/2+Y, 1/4-Z
6555	1/2+X, 1/2-Y, 3/4-Z
7555	Y, X, -Z
8555	-Y, -X, 1/2-Z

 REMARK 290
 REMARK 290 WHERE NNN -> OPERATOR NUMBER
 REMARK 290 MMM -> TRANSLATION VECTOR
 REMARK 290
 REMARK 290 CRYSTALLOGRAPHIC SYMMETRY TRANSFORMATIONS
 REMARK 290 THE FOLLOWING TRANSFORMATIONS OPERATE ON THE ATOM/HETATM
 REMARK 290 RECORDS IN THIS ENTRY TO PRODUCE CRYSTALLOGRAPHICALLY
 REMARK 290 RELATED MOLECULES.

SMTRY1	1	1.000000	0.000000	0.000000	0.000000
SMTRY2	1	0.000000	1.000000	0.000000	0.000000
SMTRY3	1	0.000000	0.000000	1.000000	0.000000
SMTRY1	2	-1.000000	0.000000	0.000000	0.000000
SMTRY2	2	0.000000	-1.000000	0.000000	0.000000
SMTRY3	2	0.000000	0.000000	1.000000	73.36600
SMTRY1	3	0.000000	-1.000000	0.000000	43.17100
SMTRY2	3	1.000000	0.000000	0.000000	43.17100
SMTRY3	3	0.000000	0.000000	1.000000	36.68300

REMARK 290 SMTRY1 4 0.000000 1.000000 0.000000 43.17100
REMARK 290 SMTRY2 4 -1.000000 0.000000 0.000000 43.17100
REMARK 290 SMTRY3 4 0.000000 0.000000 1.000000 110.04900
REMARK 290 SMTRY1 5 -1.000000 0.000000 0.000000 43.17100
REMARK 290 SMTRY2 5 0.000000 1.000000 0.000000 43.17100
REMARK 290 SMTRY3 5 0.000000 0.000000 -1.000000 36.68300
REMARK 290 SMTRY1 6 1.000000 0.000000 0.000000 43.17100
REMARK 290 SMTRY2 6 0.000000 -1.000000 0.000000 43.17100
REMARK 290 SMTRY3 6 0.000000 0.000000 -1.000000 110.04900
REMARK 290 SMTRY1 7 0.000000 1.000000 0.000000 0.00000
REMARK 290 SMTRY2 7 1.000000 0.000000 0.000000 0.00000
REMARK 290 SMTRY3 7 0.000000 0.000000 -1.000000 0.00000
REMARK 290 SMTRY1 8 0.000000 -1.000000 0.000000 0.00000
REMARK 290 SMTRY2 8 -1.000000 0.000000 0.000000 0.00000
REMARK 290 SMTRY3 8 0.000000 0.000000 -1.000000 73.36600
REMARK 290
REMARK 290 REMARK: NULL
REMARK 300
REMARK 300 BIOMOLECULE: 1
REMARK 300 THIS ENTRY CONTAINS THE CRYSTALLOGRAPHIC ASYMMETRIC UNIT
REMARK 300 WHICH CONSISTS OF 2 CHAIN(S). SEE REMARK 350 FOR
REMARK 300 INFORMATION ON GENERATING THE BIOLOGICAL MOLECULE(S).
REMARK 300
REMARK 300 QUATERNARY STRUCTURE FOR THIS ENTRY: DIMERIC
REMARK 300
REMARK 300 THE PROTEIN IS A HOMODIMER FORMED BY CHAIN A.
REMARK 300
REMARK 300 FOR THE HOMO-ASSEMBLY DESCRIBED BY REMARK 350
REMARK 300 THE DIFFERENCE IN ACCESSIBLE SURFACE AREA PER
REMARK 300 CHAIN BETWEEN THE ISOLATED CHAIN AND THAT FOR
REMARK 300 THE CHAIN IN THE COMPLEX IS 1600.4 ANGSTROM**2
REMARK 350
REMARK 350 GENERATING THE BIOMOLECULE
REMARK 350 COORDINATES FOR A COMPLETE MULTIMER REPRESENTING THE KNOWN
REMARK 350 BIOLOGICALLY SIGNIFICANT OLIGOMERIZATION STATE OF THE
REMARK 350 MOLECULE CAN BE GENERATED BY APPLYING BIOMT TRANSFORMATIONS
REMARK 350 GIVEN BELOW. BOTH NON-CRYSTALLOGRAPHIC AND
REMARK 350 CRYSTALLOGRAPHIC OPERATIONS ARE GIVEN.
REMARK 350
REMARK 350 BIOMOLECULE: 1
REMARK 350 APPLY THE FOLLOWING TO CHAINS: A
REMARK 350 BIOMT1 1 1.000000 0.000000 0.000000 0.00000
REMARK 350 BIOMT2 1 0.000000 1.000000 0.000000 0.00000
REMARK 350 BIOMT3 1 0.000000 0.000000 1.000000 0.00000
REMARK 350 BIOMT1 2 0.000000 -1.000000 0.000000 86.34200
REMARK 350 BIOMT2 2 -1.000000 0.000000 0.000000 86.34200
REMARK 350 BIOMT3 2 0.000000 0.000000 -1.000000 73.36600
REMARK 400
REMARK 400 COMPOUND
REMARK 400
REMARK 400 THE PROTEIN (CHAIN A) WAS CRYSTALLIZED IN THE PRESENCE
REMARK 400 OF A PEPTIDE FRAGMENT FROM ENDOTHELIAL PAS DOMAIN PROTEIN 1
REMARK 400 SWISS-PROT ID Q99814 (RESIDUES 846-858) BUT NONE OF THE
REMARK 400 RESIDUES CORRESPONDING TO THE PEPTIDE WERE VISIBLE IN THE
REMARK 400 ELECTRON DENSITY MAPS. IT IS POSSIBLE THAT THE PEPTIDE DID
REMARK 400 NOT BIND TO THE PROTEIN AND HENCE HAS NOT BEEN INCLUDED IN THE
REMARK 400 COMPND, SOURCE AND SEQRES RECORDS.
REMARK 400
REMARK 400 THE SEQUENCE OF THE FRAGMENT IS GIVEN BELOW.
REMARK 400
REMARK 400 VAL ASN VAL PRO VAL LEU GLY SER SER THR LEU LEU GLN

REMARK 465
REMARK 465 MISSING RESIDUES
REMARK 465 THE FOLLOWING RESIDUES WERE NOT LOCATED IN THE
REMARK 465 EXPERIMENT. (M=MODEL NUMBER; RES=RESIDUE NAME; C=CHAIN
REMARK 465 IDENTIFIER; SSSEQ=SEQUENCE NUMBER; I=INSERTION CODE.)
REMARK 465
REMARK 465 M RES C SSSEQI
REMARK 465 MET A 1
REMARK 465 ALA A 2
REMARK 465 ALA A 3
REMARK 465 THR A 4
REMARK 465 ALA A 5
REMARK 465 ALA A 6
REMARK 465 GLU A 7
REMARK 465 ALA A 8
REMARK 465 VAL A 9
REMARK 465 ALA A 10
REMARK 465 SER A 11
REMARK 465 GLY A 12
REMARK 465 SER A 13
REMARK 465 GLY A 14
REMARK 465 LYS A 304
REMARK 465 ARG A 305
REMARK 465 ILE A 306
REMARK 470
REMARK 470 MISSING ATOM
REMARK 470 THE FOLLOWING RESIDUES HAVE MISSING ATOMS (M=MODEL NUMBER;
REMARK 470 RES=RESIDUE NAME; C=CHAIN IDENTIFIER; SSEQ=SEQUENCE NUMBER;
REMARK 470 I=INSERTION CODE):
REMARK 470 M RES CSSEQI ATOMS
REMARK 470 GLU A 15 CG CD OE1 OE2
REMARK 470 GLU A 29 CG CD OE1 OE2
REMARK 470 ASN A 87 CG OD1 ND2
REMARK 470 LYS A 106 CD CE NZ
REMARK 470 LYS A 115 CG CD CE NZ
REMARK 470 ARG A 117 CG CD NE CZ NH1 NH2
REMARK 470 GLN A 133 CG CD OE1 NE2
REMARK 470 GLN A 136 CG CD OE1 NE2
REMARK 470 GLN A 137 CG CD OE1 NE2
REMARK 470 ARG A 156 CG CD NE CZ NH1 NH2
REMARK 470 LYS A 157 CD CE NZ
REMARK 470 LYS A 311 CG CD CE NZ
REMARK 500
REMARK 500 GEOMETRY AND STEREOCHEMISTRY
REMARK 500 SUBTOPIC: COVALENT BOND ANGLES
REMARK 500
REMARK 500 THE STEREOCHEMICAL PARAMETERS OF THE FOLLOWING RESIDUES
REMARK 500 HAVE VALUES WHICH DEVIATE FROM EXPECTED VALUES BY MORE
REMARK 500 THAN 6*RMSD (M=MODEL NUMBER; RES=RESIDUE NAME; C=CHAIN
REMARK 500 IDENTIFIER; SSEQ=SEQUENCE NUMBER; I=INSERTION CODE).
REMARK 500
REMARK 500 STANDARD TABLE:
REMARK 500 FORMAT: (10X,I3,1X,A3,1X,A1,I4,A1,3(1X,A4,2X),12X,F5.1)
REMARK 500
REMARK 500 EXPECTED VALUES: ENGH AND HUBER, 1991
REMARK 500
REMARK 500 M RES CSSEQI ATM1 ATM2 ATM3
REMARK 500 TYR A 261 N - CA - C ANGL. DEV. = -10.1 DEGREES
REMARK 500
REMARK 500 REMARK: NULL
REMARK 500

REMARK 500 GEOMETRY AND STEREOCHEMISTRY
 REMARK 500 SUBTOPIC: COVALENT BOND LENGTHS
 REMARK 500
 REMARK 500 THE STEREOCHEMICAL PARAMETERS OF THE FOLLOWING RESIDUES
 REMARK 500 HAVE VALUES WHICH DEVIATE FROM EXPECTED VALUES BY MORE
 REMARK 500 THAN 6*RMSD AND BY MORE THAN 0.150 ANGSTROMS (M=MODEL
 REMARK 500 NUMBER; RES=RESIDUE NAME; C=CHAIN IDENTIFIER; SSEQ=SEQUENCE
 REMARK 500 NUMBER; I=INSERTION CODE).
 REMARK 500
 REMARK 500 STANDARD TABLE:
 REMARK 500 FORMAT: (10X,I3,1X,A3,1X,A1,I4,A1,1X,2(A4,A1,3X),12X,F5.3)
 REMARK 500
 REMARK 500 EXPECTED VALUESS: ENGH AND HUBER, 1991
 REMARK 500

REMARK 500	M	RES	CSSEQI	ATM1	RES	CSSEQI	ATM2	DEVIATION
REMARK 500		MET	A	343	SD	MET	A 343	CE
REMARK 500								-0.151

 REMARK 500
 REMARK 500 REMARK: NULL
 REMARK 500
 REMARK 500 GEOMETRY AND STEREOCHEMISTRY
 REMARK 500 SUBTOPIC: CLOSE CONTACTS IN SAME ASYMMETRIC UNIT
 REMARK 500
 REMARK 500 THE FOLLOWING ATOMS ARE IN CLOSE CONTACT.
 REMARK 500

REMARK 500	ATM1	RES	C	SSEQI	ATM2	RES	C	SSEQI	DISTANCE
REMARK 500	OG1	THR	A	39	OE1	GLU	A	262	2.16

 REMARK 525
 REMARK 525 SOLVENT
 REMARK 525
 REMARK 525 THE SOLVENT MOLECULES ARE GIVEN CHAIN IDENTIFIERS TO
 REMARK 525 INDICATE THE PROTEIN CHAIN TO WHICH THEY ARE MOST CLOSELY
 REMARK 525 ASSOCIATED WITH:

REMARK 525	PROTEIN CHAIN	SOLVENT CHAIN
REMARK 525	A	Z

 REMARK 600
 REMARK 600 HETEROGEN
 REMARK 600
 REMARK 600 FOR METAL ATOM FE FE2 A1350 THE COORDINATION ANGLES ARE:

REMARK 600	1	HIS	199A	NE2				
REMARK 600	2	ASP	201A	OD2	106.1			
REMARK 600	3	HIS	279A	NE2	77.8	85.5		
REMARK 600	4	AKG	1351A	O1	168.0	81.1	93.4	
REMARK 600	5	AKG	1351A	O5	88.7	163.3	90.2	83.1
REMARK 600					1	2	3	4

 REMARK 700
 REMARK 700 SHEET
 REMARK 700 THE SHEET STRUCTURE OF THIS MOLECULE IS BIFURCATED. IN
 REMARK 700 ORDER TO REPRESENT THIS FEATURE IN THE SHEET RECORDS BELOW,
 REMARK 700 TWO SHEETS ARE DEFINED.
 REMARK 800
 REMARK 800 SITE
 REMARK 800 SITE_IDENTIFIER: FEA
 REMARK 800 SITE_DESCRIPTION: FE BINDING SITE FOR CHAIN A
 REMARK 800
 REMARK 800 SITE_IDENTIFIER: AKG
 REMARK 800 SITE_DESCRIPTION: AKG BINDING SITE FOR CHAIN A
 REMARK 800
 REMARK 800 SITE_IDENTIFIER: SA1
 REMARK 800 SITE_DESCRIPTION: SO4 BINDING SITE FOR CHAIN A
 REMARK 800

REMARK 800 SITE_IDENTIFIER: SA2
 REMARK 800 SITE_DESCRIPTION: SO4 BINDING SITE FOR CHAIN A
 REMARK 800
 REMARK 800 SITE_IDENTIFIER: SA3
 REMARK 800 SITE_DESCRIPTION: SO4 BINDING SITE FOR CHAIN A
 REMARK 900
 REMARK 900 RELATED ENTRIES
 REMARK 900 RELATED ID: 1H2K RELATED DB: PDB
 REMARK 900 FACTOR INHIBITING HIF-1 ALPHA IN COMPLEX
 REMARK 900 WITH HIF-1 ALPHA FRAGMENT PEPTIDE
 REMARK 900 RELATED ID: 1H2L RELATED DB: PDB
 REMARK 900 FACTOR INHIBITING HIF-1 ALPHA IN COMPLEX
 REMARK 900 WITH HIF-1 ALPHA FRAGMENT PEPTIDE
 REMARK 900 RELATED ID: 1H2M RELATED DB: PDB
 REMARK 900 FACTOR INHIBITING HIF-1 ALPHA IN COMPLEX
 REMARK 900 WITH HIF-1 ALPHA FRAGMENT PEPTIDE

DBREF 1H2N A 1 349 SWS Q969Q7 Q969Q7 1 349
 SEQRES 1 A 349 MET ALA ALA THR ALA ALA GLU ALA VAL ALA SER GLY SER
 SEQRES 2 A 349 GLY GLU PRO ARG GLU GLU ALA GLY ALA LEU GLY PRO ALA
 SEQRES 3 A 349 TRP ASP GLU SER GLN LEU ARG SER TYR SER PHE PRO THR
 SEQRES 4 A 349 ARG PRO ILE PRO ARG LEU SER GLN SER ASP PRO ARG ALA
 SEQRES 5 A 349 GLU GLU LEU ILE GLU ASN GLU GLU PRO VAL VAL LEU THR
 SEQRES 6 A 349 ASP THR ASN LEU VAL TYR PRO ALA LEU LYS TRP ASP LEU
 SEQRES 7 A 349 GLU TYR LEU GLN GLU ASN ILE GLY ASN GLY ASP PHE SER
 SEQRES 8 A 349 VAL TYR SER ALA SER THR HIS LYS PHE LEU TYR TYR ASP
 SEQRES 9 A 349 GLU LYS LYS MET ALA ASN PHE GLN ASN PHE LYS PRO ARG
 SEQRES 10 A 349 SER ASN ARG GLU GLU MET LYS PHE HIS GLU PHE VAL GLU
 SEQRES 11 A 349 LYS LEU GLN ASP ILE GLN GLN ARG GLY GLY GLU GLU ARG
 SEQRES 12 A 349 LEU TYR LEU GLN GLN THR LEU ASN ASP THR VAL GLY ARG
 SEQRES 13 A 349 LYS ILE VAL MET ASP PHE LEU GLY PHE ASN TRP ASN TRP
 SEQRES 14 A 349 ILE ASN LYS GLN GLN GLY LYS ARG GLY TRP GLY GLN LEU
 SEQRES 15 A 349 THR SER ASN LEU LEU LEU ILE GLY MET GLU GLY ASN VAL
 SEQRES 16 A 349 THR PRO ALA HIS TYR ASP GLU GLN GLN ASN PHE PHE ALA
 SEQRES 17 A 349 GLN ILE LYS GLY TYR LYS ARG CYS ILE LEU PHE PRO PRO
 SEQRES 18 A 349 ASP GLN PHE GLU CYS LEU TYR PRO TYR PRO VAL HIS HIS
 SEQRES 19 A 349 PRO CYS ASP ARG GLN SER GLN VAL ASP PHE ASP ASN PRO
 SEQRES 20 A 349 ASP TYR GLU ARG PHE PRO ASN PHE GLN ASN VAL VAL GLY
 SEQRES 21 A 349 TYR GLU THR VAL VAL GLY PRO GLY ASP VAL LEU TYR ILE
 SEQRES 22 A 349 PRO MET TYR TRP TRP HIS HIS ILE GLU SER LEU LEU ASN
 SEQRES 23 A 349 GLY GLY ILE THR ILE THR VAL ASN PHE TRP TYR LYS GLY
 SEQRES 24 A 349 ALA PRO THR PRO LYS ARG ILE GLU TYR PRO LEU LYS ALA
 SEQRES 25 A 349 HIS GLN LYS VAL ALA ILE MET ARG ASN ILE GLU LYS MET
 SEQRES 26 A 349 LEU GLY GLU ALA LEU GLY ASN PRO GLN GLU VAL GLY PRO
 SEQRES 27 A 349 LEU LEU ASN THR MET ILE LYS GLY ARG TYR ASN

HET FE2 A1350 1
 HET AKG A1351 10
 HET SO4 A1352 5
 HET SO4 A1353 5
 HET SO4 A1354 5
 HETNAM FE2 FE (II) ION
 HETNAM AKG 2-OXYGLUTARIC ACID
 HETNAM SO4 SULFATE ION
 FORMUL 3 FE2 FE1 2+
 FORMUL 4 AKG C5 H6 O5
 FORMUL 5 SO4 3(O4 S1 2-)
 FORMUL 6 HOH *3(H2 O1)

HELIX 1 1 ASP A 28 LEU A 32 5
 HELIX 2 2 ASP A 49 ASN A 58 1
 HELIX 3 3 VAL A 70 TRP A 76 5
 HELIX 4 4 ASP A 77 ILE A 85 1
 HELIX 5 5 ASP A 104 GLN A 112 5

5
 10
 7
 9
 9

[illegible]

ATOM	11	CG	PRO	A	16	4.749	30.155	12.612	1.00	42.41	C
ATOM	12	CD	PRO	A	16	5.103	30.916	11.382	1.00	42.22	C
ATOM	13	N	ARG	A	17	7.231	26.791	10.446	1.00	42.51	N
ATOM	14	CA	ARG	A	17	7.337	25.723	9.478	1.00	42.62	C
ATOM	15	C	ARG	A	17	6.095	24.856	9.527	1.00	42.85	C
ATOM	16	O	ARG	A	17	5.492	24.705	10.587	1.00	43.51	O
ATOM	17	CB	ARG	A	17	8.505	24.800	9.845	1.00	42.50	C
ATOM	18	CG	ARG	A	17	9.871	25.434	9.859	1.00	42.22	C
ATOM	19	CD	ARG	A	17	10.995	24.466	10.228	1.00	42.12	C
ATOM	20	NE	ARG	A	17	11.085	24.138	11.656	1.00	42.49	N
ATOM	21	CZ	ARG	A	17	11.588	24.943	12.607	1.00	42.43	C
ATOM	22	NH1	ARG	A	17	12.033	26.158	12.316	1.00	42.11	N
ATOM	23	NH2	ARG	A	17	11.649	24.529	13.869	1.00	42.25	N
ATOM	24	N	GLU	A	18	5.742	24.242	8.404	1.00	42.60	N
ATOM	25	CA	GLU	A	18	4.662	23.268	8.405	1.00	42.36	C
ATOM	26	C	GLU	A	18	5.203	21.870	8.728	1.00	41.90	C
ATOM	27	O	GLU	A	18	6.249	21.472	8.216	1.00	41.98	O
ATOM	28	CB	GLU	A	18	3.988	23.224	7.041	1.00	42.57	C
ATOM	29	CG	GLU	A	18	3.363	24.537	6.620	1.00	43.60	C
ATOM	30	CD	GLU	A	18	1.993	24.756	7.233	1.00	45.11	C
ATOM	31	OE1	GLU	A	18	1.365	23.766	7.686	1.00	45.48	O
ATOM	32	OE2	GLU	A	18	1.546	25.926	7.253	1.00	46.49	O
ATOM	33	N	GLU	A	19	4.487	21.133	9.578	1.00	41.27	N
ATOM	34	CA	GLU	A	19	4.806	19.737	9.894	1.00	40.61	C
ATOM	35	C	GLU	A	19	4.478	18.802	8.748	1.00	39.73	C
ATOM	36	O	GLU	A	19	3.424	18.915	8.137	1.00	39.76	O
ATOM	37	CB	GLU	A	19	4.024	19.299	11.128	1.00	40.72	C
ATOM	38	CG	GLU	A	19	4.507	20.034	12.361	1.00	42.24	C
ATOM	39	CD	GLU	A	19	4.358	19.278	13.668	1.00	44.37	C
ATOM	40	OE1	GLU	A	19	3.852	18.127	13.680	1.00	46.17	O
ATOM	41	OE2	GLU	A	19	4.769	19.863	14.700	1.00	45.44	O
ATOM	42	N	ALA	A	20	5.369	17.859	8.478	1.00	38.99	N
ATOM	43	CA	ALA	A	20	5.164	16.922	7.390	1.00	38.71	C
ATOM	44	C	ALA	A	20	3.831	16.232	7.542	1.00	38.51	C
ATOM	45	O	ALA	A	20	3.391	15.933	8.649	1.00	38.80	O
ATOM	46	CB	ALA	A	20	6.280	15.895	7.328	1.00	38.51	C
ATOM	47	N	GLY	A	21	3.180	15.991	6.419	1.00	38.03	N
ATOM	48	CA	GLY	A	21	1.924	15.290	6.449	1.00	38.11	C
ATOM	49	C	GLY	A	21	0.746	16.205	6.682	1.00	38.19	C
ATOM	50	O	GLY	A	21	-0.328	15.743	7.049	1.00	38.25	O
ATOM	51	N	ALA	A	22	0.941	17.497	6.447	1.00	38.40	N
ATOM	52	CA	ALA	A	22	-0.130	18.472	6.571	1.00	38.50	C
ATOM	53	C	ALA	A	22	-0.725	18.413	7.960	1.00	38.60	C
ATOM	54	O	ALA	A	22	-1.930	18.547	8.142	1.00	38.65	O
ATOM	55	CB	ALA	A	22	-1.196	18.242	5.520	1.00	38.61	C
ATOM	56	N	LEU	A	23	0.135	18.206	8.946	1.00	38.75	N
ATOM	57	CA	LEU	A	23	-0.297	18.257	10.330	1.00	38.82	C
ATOM	58	C	LEU	A	23	-0.321	19.684	10.843	1.00	38.78	C
ATOM	59	O	LEU	A	23	-0.525	19.919	12.030	1.00	38.74	O
ATOM	60	CB	LEU	A	23	0.597	17.392	11.193	1.00	38.77	C
ATOM	61	CG	LEU	A	23	0.421	15.952	10.736	1.00	39.03	C
ATOM	62	CD1	LEU	A	23	1.203	14.958	11.579	1.00	39.08	C
ATOM	63	CD2	LEU	A	23	-1.070	15.636	10.750	1.00	39.43	C
ATOM	64	N	GLY	A	24	-0.094	20.636	9.947	1.00	38.75	N
ATOM	65	CA	GLY	A	24	-0.248	22.025	10.298	1.00	38.79	C
ATOM	66	C	GLY	A	24	0.972	22.669	10.861	1.00	38.90	C
ATOM	67	O	GLY	A	24	2.042	22.084	10.961	1.00	39.06	O
ATOM	68	N	PRO	A	25	0.791	23.906	11.267	1.00	39.19	N
ATOM	69	CA	PRO	A	25	1.903	24.682	11.778	1.00	39.35	C
ATOM	70	C	PRO	A	25	2.332	24.037	13.074	1.00	39.62	C
ATOM	71	O	PRO	A	25	1.492	23.633	13.871	1.00	39.48	O

ATOM	72	CB	PRO	A	25	1.299	26.062	12.019	1.00	39.24	
ATOM	73	CG	PRO	A	25	-0.142	25.920	11.961	1.00	38.71	C
ATOM	74	CD	PRO	A	25	-0.490	24.616	11.365	1.00	39.17	C
ATOM	75	N	ALA	A	26	3.631	23.899	13.252	1.00	40.22	N
ATOM	76	CA	ALA	A	26	4.170	23.342	14.476	1.00	40.79	C
ATOM	77	C	ALA	A	26	3.930	24.294	15.666	1.00	41.07	C
ATOM	78	O	ALA	A	26	3.769	23.837	16.797	1.00	41.57	O
ATOM	79	CB	ALA	A	26	5.633	23.068	14.300	1.00	40.98	C
ATOM	80	N	TRP	A	27	3.931	25.605	15.429	1.00	40.80	N
ATOM	81	CA	TRP	A	27	3.543	26.552	16.477	1.00	40.67	C
ATOM	82	C	TRP	A	27	2.982	27.848	15.880	1.00	40.33	C
ATOM	83	O	TRP	A	27	2.777	27.940	14.675	1.00	40.13	O
ATOM	84	CB	TRP	A	27	4.742	26.860	17.358	1.00	40.70	C
ATOM	85	CG	TRP	A	27	5.942	27.014	16.558	1.00	41.36	C
ATOM	86	CD1	TRP	A	27	6.762	26.023	16.074	1.00	43.03	C
ATOM	87	CD2	TRP	A	27	6.466	28.233	16.084	1.00	40.41	C
ATOM	88	NE1	TRP	A	27	7.788	26.579	15.344	1.00	42.65	N
ATOM	89	CE2	TRP	A	27	7.618	27.938	15.331	1.00	41.20	C
ATOM	90	CE3	TRP	A	27	6.081	29.555	16.223	1.00	40.47	C
ATOM	91	CZ2	TRP	A	27	8.370	28.912	14.732	1.00	41.10	C
ATOM	92	CZ3	TRP	A	27	6.827	30.514	15.640	1.00	41.11	C
ATOM	93	CH2	TRP	A	27	7.962	30.198	14.900	1.00	41.57	C
ATOM	94	N	ASP	A	28	2.677	28.830	16.723	1.00	39.99	N
ATOM	95	CA	ASP	A	28	2.322	30.143	16.209	1.00	39.78	C
ATOM	96	C	ASP	A	28	2.802	31.257	17.121	1.00	39.17	C
ATOM	97	O	ASP	A	28	3.227	31.013	18.240	1.00	39.00	O
ATOM	98	CB	ASP	A	28	0.826	30.254	15.981	1.00	40.16	C
ATOM	99	CG	ASP	A	28	0.066	30.380	17.258	1.00	40.79	C
ATOM	100	OD1	ASP	A	28	0.044	31.496	17.821	1.00	41.30	O
ATOM	101	OD2	ASP	A	28	-0.531	29.414	17.774	1.00	42.25	O
ATOM	102	N	GLU	A	29	2.720	32.486	16.624	1.00	38.73	N
ATOM	103	CA	GLU	A	29	3.223	33.663	17.341	1.00	38.23	C
ATOM	104	C	GLU	A	29	2.739	33.764	18.781	1.00	37.69	C
ATOM	105	O	GLU	A	29	3.492	34.131	19.664	1.00	37.34	O
ATOM	106	CB	GLU	A	29	2.834	34.924	16.594	1.00	38.20	C
ATOM	107	N	SER	A	30	1.482	33.418	19.012	1.00	37.46	N
ATOM	108	CA	SER	A	30	0.874	33.583	20.324	1.00	37.18	C
ATOM	109	C	SER	A	30	1.562	32.774	21.399	1.00	36.93	C
ATOM	110	O	SER	A	30	1.282	32.949	22.577	1.00	36.82	O
ATOM	111	CB	SER	A	30	-0.595	33.164	20.284	1.00	37.23	C
ATOM	112	OG	SER	A	30	-0.744	31.792	20.619	1.00	36.97	O
ATOM	113	N	GLN	A	31	2.441	31.867	20.999	1.00	36.85	N
ATOM	114	CA	GLN	A	31	3.128	31.021	21.961	1.00	36.77	C
ATOM	115	C	GLN	A	31	4.445	31.636	22.340	1.00	36.59	C
ATOM	116	O	GLN	A	31	5.141	31.127	23.220	1.00	36.74	O
ATOM	117	CB	GLN	A	31	3.366	29.621	21.395	1.00	36.71	C
ATOM	118	CG	GLN	A	31	2.084	28.828	21.234	1.00	36.97	C
ATOM	119	CD	GLN	A	31	2.282	27.497	20.560	1.00	36.68	C
ATOM	120	OE1	GLN	A	31	2.133	27.386	19.346	1.00	36.62	O
ATOM	121	NE2	GLN	A	31	2.601	26.478	21.343	1.00	36.85	N
ATOM	122	N	LEU	A	32	4.794	32.726	21.670	1.00	36.32	N
ATOM	123	CA	LEU	A	32	6.050	33.381	21.942	1.00	36.17	C
ATOM	124	C	LEU	A	32	5.817	34.498	22.921	1.00	36.09	C
ATOM	125	O	LEU	A	32	4.837	35.233	22.815	1.00	36.18	O
ATOM	126	CB	LEU	A	32	6.673	33.928	20.664	1.00	36.07	C
ATOM	127	CG	LEU	A	32	6.990	32.871	19.604	1.00	36.72	C
ATOM	128	CD1	LEU	A	32	7.747	33.457	18.453	1.00	36.96	C
ATOM	129	CD2	LEU	A	32	7.797	31.744	20.161	1.00	37.31	C
ATOM	130	N	ARG	A	33	6.700	34.576	23.908	1.00	35.99	N
ATOM	131	CA	ARG	A	33	6.713	35.671	24.851	1.00	36.15	C
ATOM	132	C	ARG	A	33	7.171	36.941	24.130	1.00	35.95	C

ATOM	133	O	ARG	A	33	7.950	36.888	23.188	1.00	35.75
ATOM	134	CB	ARG	A	33	7.671	35.350	25.991	1.00	36.41
ATOM	135	CG	ARG	A	33	7.210	34.204	26.854	1.00	36.76
ATOM	136	CD	ARG	A	33	8.082	33.948	28.070	1.00	36.81
ATOM	137	NE	ARG	A	33	7.479	32.927	28.923	1.00	37.39
ATOM	138	CZ	ARG	A	33	6.501	33.160	29.785	1.00	37.61
ATOM	139	NH1	ARG	A	33	6.027	34.385	29.931	1.00	37.75
ATOM	140	NH2	ARG	A	33	6.001	32.171	30.513	1.00	38.47
ATOM	141	N	SER	A	34	6.707	38.088	24.588	1.00	35.92
ATOM	142	CA	SER	A	34	7.017	39.331	23.902	1.00	36.05
ATOM	143	C	SER	A	34	8.044	40.146	24.671	1.00	35.4
ATOM	144	O	SER	A	34	7.940	40.311	25.884	1.00	35.41
ATOM	145	CB	SER	A	34	5.732	40.120	23.691	1.00	36.27
ATOM	146	OG	SER	A	34	5.046	40.253	24.920	1.00	37.99
ATOM	147	N	TYR	A	35	9.031	40.655	23.945	1.00	35.01
ATOM	148	CA	TYR	A	35	10.140	41.370	24.543	1.00	34.69
ATOM	149	C	TYR	A	35	10.400	42.693	23.853	1.00	34.59
ATOM	150	O	TYR	A	35	9.841	42.989	22.813	1.00	34.45
ATOM	151	CB	TYR	A	35	11.384	40.501	24.469	1.00	34.71
ATOM	152	CG	TYR	A	35	11.228	39.222	25.232	1.00	34.43
ATOM	153	CD1	TYR	A	35	10.928	39.246	26.573	1.00	33.87
ATOM	154	CD2	TYR	A	35	11.350	37.986	24.602	1.00	35.32
ATOM	155	CE1	TYR	A	35	10.775	38.087	27.285	1.00	34.96
ATOM	156	CE2	TYR	A	35	11.192	36.807	25.309	1.00	35.50
ATOM	157	CZ	TYR	A	35	10.904	36.865	26.653	1.00	35.16
ATOM	158	OH	TYR	A	35	10.742	35.709	27.376	1.00	34.77
ATOM	159	N	SER	A	36	11.290	43.481	24.429	1.00	34.77
ATOM	160	CA	SER	A	36	11.572	44.814	23.924	1.00	34.74
ATOM	161	C	SER	A	36	12.567	44.926	22.771	1.00	34.30
ATOM	162	O	SER	A	36	12.805	46.011	22.287	1.00	34.23
ATOM	163	CB	SER	A	36	12.127	45.627	25.077	1.00	34.91
ATOM	164	OG	SER	A	36	13.395	45.116	25.449	1.00	35.56
ATOM	165	N	PHE	A	37	13.155	43.838	22.318	1.00	34.15
ATOM	166	CA	PHE	A	37	14.193	43.963	21.306	1.00	34.11
ATOM	167	C	PHE	A	37	13.969	43.177	20.044	1.00	34.07
ATOM	168	O	PHE	A	37	13.356	42.109	20.017	1.00	33.98
ATOM	169	CB	PHE	A	37	15.514	43.479	21.867	1.00	34.08
ATOM	170	CG	PHE	A	37	15.417	42.147	22.438	1.00	34.33
ATOM	171	CD1	PHE	A	37	15.445	41.034	21.621	1.00	35.66
ATOM	172	CD2	PHE	A	37	15.208	41.990	23.777	1.00	35.42
ATOM	173	CE1	PHE	A	37	15.320	39.782	22.143	1.00	35.73
ATOM	174	CE2	PHE	A	37	15.084	40.736	24.317	1.00	36.14
ATOM	175	CZ	PHE	A	37	15.141	39.628	23.497	1.00	36.55
ATOM	176	N	PRO	A	38	14.565	43.682	18.991	1.00	33.95
ATOM	177	CA	PRO	A	38	14.474	43.028	17.704	1.00	33.91
ATOM	178	C	PRO	A	38	15.445	41.878	17.640	1.00	33.74
ATOM	179	O	PRO	A	38	16.366	41.803	18.441	1.00	33.82
ATOM	180	CB	PRO	A	38	14.887	44.133	16.751	1.00	34.03
ATOM	181	CG	PRO	A	38	15.874	44.935	17.548	1.00	33.99
ATOM	182	CD	PRO	A	38	15.385	44.903	18.945	1.00	33.80
ATOM	183	N	THR	A	39	15.233	40.996	16.678	1.00	34.02
ATOM	184	CA	THR	A	39	16.089	39.844	16.464	1.00	34.00
ATOM	185	C	THR	A	39	16.035	39.490	15.008	1.00	34.09
ATOM	186	O	THR	A	39	15.137	39.894	14.311	1.00	34.05
ATOM	187	CB	THR	A	39	15.563	38.637	17.201	1.00	33.94
ATOM	188	OG1	THR	A	39	14.250	38.336	16.714	1.00	34.18
ATOM	189	CG2	THR	A	39	15.338	38.915	18.642	1.00	34.09
ATOM	190	N	ARG	A	40	16.992	38.695	14.576	1.00	34.62
ATOM	191	CA	ARG	A	40	17.066	38.204	13.228	1.00	35.13
ATOM	192	C	ARG	A	40	17.100	36.721	13.361	1.00	35.17
ATOM	193	O	ARG	A	40	17.419	36.189	14.434	1.00	34.96

[illegible]

ATOM	194	CB	ARG	A	40	18.352	38.649	12.568	1.00	35.54
ATOM	195	CG	ARG	A	40	18.358	40.103	12.287	1.00	38.87
ATOM	196	CD	ARG	A	40	16.975	40.597	11.893	1.00	42.88
ATOM	197	NE	ARG	A	40	16.737	40.908	10.489	1.00	44.50
ATOM	198	CZ	ARG	A	40	15.554	41.298	10.076	1.00	46.73
ATOM	199	NH1	ARG	A	40	14.574	41.350	10.974	1.00	46.73
ATOM	200	NH2	ARG	A	40	15.334	41.636	8.802	1.00	48.87
ATOM	201	N	PRO	A	41	16.775	36.026	12.285	1.00	35.20
ATOM	202	CA	PRO	A	41	16.749	34.579	12.348	1.00	35.24
ATOM	203	C	PRO	A	41	18.114	33.973	12.275	1.00	35.10
ATOM	204	O	PRO	A	41	18.995	34.455	11.573	1.00	35.96
ATOM	205	CB	PRO	A	41	16.003	34.208	11.079	1.00	35.27
ATOM	206	CG	PRO	A	41	15.408	35.453	10.640	1.00	34.93
ATOM	207	CD	PRO	A	41	16.381	36.494	10.953	1.00	34.81
ATOM	208	N	ILE	A	42	18.277	32.910	13.022	1.00	34.55
ATOM	209	CA	ILE	A	42	19.435	32.099	12.909	1.00	34.29
ATOM	210	C	ILE	A	42	19.145	31.194	11.713	1.00	34.29
ATOM	211	O	ILE	A	42	18.035	30.654	11.598	1.00	34.38
ATOM	212	CB	ILE	A	42	19.545	31.258	14.150	1.00	34.14
ATOM	213	CG1	ILE	A	42	19.742	32.152	15.368	1.00	34.83
ATOM	214	CG2	ILE	A	42	20.679	30.297	14.016	1.00	34.41
ATOM	215	CD1	ILE	A	42	19.479	31.467	16.678	1.00	35.37
ATOM	216	N	PRO	A	43	20.127	31.010	10.836	1.00	33.80
ATOM	217	CA	PRO	A	43	19.982	30.122	9.686	1.00	33.62
ATOM	218	C	PRO	A	43	19.714	28.676	10.066	1.00	33.65
ATOM	219	O	PRO	A	43	20.320	28.198	11.015	1.00	33.60
ATOM	220	CB	PRO	A	43	21.366	30.174	9.043	1.00	33.63
ATOM	221	CG	PRO	A	43	21.986	31.385	9.543	1.00	33.11
ATOM	222	CD	PRO	A	43	21.455	31.630	10.882	1.00	33.51
ATOM	223	N	ARG	A	44	18.820	28.007	9.341	1.00	33.86
ATOM	224	CA	ARG	A	44	18.597	26.572	9.486	1.00	34.00
ATOM	225	C	ARG	A	44	19.176	25.968	8.247	1.00	33.87
ATOM	226	O	ARG	A	44	18.783	26.319	7.152	1.00	33.76
ATOM	227	CB	ARG	A	44	17.125	26.185	9.520	1.00	34.10
ATOM	228	CG	ARG	A	44	16.301	26.879	10.577	1.00	35.44
ATOM	229	CD	ARG	A	44	14.781	26.505	10.583	1.00	37.36
ATOM	230	NE	ARG	A	44	14.415	25.108	10.265	1.00	37.59
ATOM	231	CZ	ARG	A	44	14.237	24.144	11.187	1.00	39.71
ATOM	232	NH1	ARG	A	44	14.429	24.396	12.487	1.00	39.94
ATOM	233	NH2	ARG	A	44	13.876	22.915	10.821	1.00	40.07
ATOM	234	N	LEU	A	45	20.086	25.030	8.407	1.00	33.98
ATOM	235	CA	LEU	A	45	20.747	24.467	7.269	1.00	34.15
ATOM	236	C	LEU	A	45	20.950	22.995	7.389	1.00	34.37
ATOM	237	O	LEU	A	45	20.901	22.453	8.486	1.00	34.52
ATOM	238	CB	LEU	A	45	22.116	25.090	7.183	1.00	34.25
ATOM	239	CG	LEU	A	45	22.080	26.576	6.905	1.00	34.80
ATOM	240	CD1	LEU	A	45	23.454	27.095	7.177	1.00	36.05
ATOM	241	CD2	LEU	A	45	21.655	26.837	5.481	1.00	34.78
ATOM	242	N	SER	A	46	21.199	22.343	6.255	1.00	34.55
ATOM	243	CA	SER	A	46	21.549	20.949	6.306	1.00	34.60
ATOM	244	C	SER	A	46	22.985	20.838	6.681	1.00	34.89
ATOM	245	O	SER	A	46	23.826	21.671	6.383	1.00	34.54
ATOM	246	CB	SER	A	46	21.356	20.206	5.006	1.00	34.64
ATOM	247	OG	SER	A	46	22.231	19.090	4.989	1.00	33.61
ATOM	248	N	GLN	A	47	23.247	19.736	7.324	1.00	35.61
ATOM	249	CA	GLN	A	47	24.539	19.462	7.866	1.00	36.10
ATOM	250	C	GLN	A	47	25.565	19.368	6.746	1.00	35.96
ATOM	251	O	GLN	A	47	26.754	19.537	6.969	1.00	36.13
ATOM	252	CB	GLN	A	47	24.399	18.168	8.669	1.00	36.22
ATOM	253	CG	GLN	A	47	25.604	17.340	8.730	1.00	37.25
ATOM	254	CD	GLN	A	47	25.724	16.459	7.532	1.00	39.03

C
C
C
N
C
N
N
N
C
C
O
C
C
C
N
C
C
O
C
C
C
N
C
C
O
C
C
C
N
C
N
N
N
C
C
C
O
C
C
C
N
C
C
O
C
O
N
C
C
C
C
C

ATOM	255	OE1	GLN	A	47	24.766	16.306	6.759	1.00	38.79	O
ATOM	256	NE2	GLN	A	47	26.902	15.872	7.354	1.00	41.44	N
ATOM	257	N	SER	A	48	25.088	19.146	5.531	1.00	35.87	N
ATOM	258	CA	SER	A	48	25.958	18.938	4.391	1.00	35.90	C
ATOM	259	C	SER	A	48	26.301	20.244	3.715	1.00	36.06	C
ATOM	260	O	SER	A	48	27.215	20.334	2.897	1.00	35.95	O
ATOM	261	CB	SER	A	48	25.214	18.073	3.402	1.00	35.97	C
ATOM	262	OG	SER	A	48	23.933	18.635	3.158	1.00	36.62	O
ATOM	263	N	ASP	A	49	25.552	21.266	4.075	1.00	36.34	N
ATOM	264	CA	ASP	A	49	25.701	22.564	3.491	1.00	36.65	C
ATOM	265	C	ASP	A	49	26.963	23.236	3.952	1.00	37.06	C
ATOM	266	O	ASP	A	49	27.114	23.507	5.138	1.00	37.24	O
ATOM	267	CB	ASP	A	49	24.545	23.405	3.953	1.00	36.80	C
ATOM	268	CG	ASP	A	49	24.441	24.674	3.206	1.00	37.08	C
ATOM	269	OD1	ASP	A	49	25.490	25.217	2.810	1.00	37.38	O
ATOM	270	OD2	ASP	A	49	23.343	25.187	2.953	1.00	38.64	O
ATOM	271	N	PRO	A	50	27.857	23.565	3.029	1.00	37.42	N
ATOM	272	CA	PRO	A	50	29.113	24.196	3.430	1.00	37.32	C
ATOM	273	C	PRO	A	50	28.881	25.413	4.298	1.00	37.46	C
ATOM	274	O	PRO	A	50	29.742	25.727	5.113	1.00	37.85	O
ATOM	275	CB	PRO	A	50	29.750	24.604	2.108	1.00	37.21	C
ATOM	276	CG	PRO	A	50	29.134	23.726	1.086	1.00	37.39	C
ATOM	277	CD	PRO	A	50	27.755	23.395	1.568	1.00	37.34	C
ATOM	278	N	ARG	A	51	27.743	26.081	4.162	1.00	37.64	N
ATOM	279	CA	ARG	A	51	27.535	27.301	4.932	1.00	38.06	C
ATOM	280	C	ARG	A	51	27.470	27.000	6.421	1.00	38.26	C
ATOM	281	O	ARG	A	51	27.920	27.798	7.250	1.00	38.47	O
ATOM	282	CB	ARG	A	51	26.282	28.055	4.470	1.00	38.14	C
ATOM	283	CG	ARG	A	51	26.418	28.692	3.059	1.00	38.89	C
ATOM	284	CD	ARG	A	51	25.117	29.259	2.455	1.00	39.65	C
ATOM	285	NE	ARG	A	51	24.121	28.211	2.175	1.00	41.25	N
ATOM	286	CZ	ARG	A	51	22.798	28.404	2.078	1.00	41.86	C
ATOM	287	NH1	ARG	A	51	22.264	29.605	2.225	1.00	43.70	N
ATOM	288	NH2	ARG	A	51	21.994	27.393	1.818	1.00	41.90	N
ATOM	289	N	ALA	A	52	26.936	25.837	6.770	1.00	38.26	N
ATOM	290	CA	ALA	A	52	26.807	25.499	8.171	1.00	38.30	C
ATOM	291	C	ALA	A	52	28.206	25.374	8.683	1.00	38.45	C
ATOM	292	O	ALA	A	52	28.572	25.922	9.732	1.00	38.37	O
ATOM	293	CB	ALA	A	52	26.085	24.208	8.344	1.00	38.25	C
ATOM	294	N	GLU	A	53	29.003	24.665	7.901	1.00	38.56	N
ATOM	295	CA	GLU	A	53	30.361	24.420	8.291	1.00	38.94	C
ATOM	296	C	GLU	A	53	31.145	25.717	8.484	1.00	39.00	C
ATOM	297	O	GLU	A	53	31.970	25.793	9.386	1.00	38.49	O
ATOM	298	CB	GLU	A	53	31.060	23.507	7.314	1.00	38.91	C
ATOM	299	CG	GLU	A	53	32.139	22.738	8.035	1.00	40.55	C
ATOM	300	CD	GLU	A	53	31.662	21.403	8.579	1.00	42.38	C
ATOM	301	OE1	GLU	A	53	30.475	21.274	8.957	1.00	43.42	O
ATOM	302	OE2	GLU	A	53	32.498	20.478	8.639	1.00	44.15	O
ATOM	303	N	GLU	A	54	30.897	26.737	7.662	1.00	39.21	N
ATOM	304	CA	GLU	A	54	31.592	28.004	7.870	1.00	39.44	C
ATOM	305	C	GLU	A	54	31.092	28.625	9.144	1.00	39.05	C
ATOM	306	O	GLU	A	54	31.848	29.262	9.869	1.00	39.51	O
ATOM	307	CB	GLU	A	54	31.337	29.023	6.772	1.00	39.59	C
ATOM	308	CG	GLU	A	54	31.940	28.693	5.433	1.00	41.75	C
ATOM	309	CD	GLU	A	54	31.098	29.263	4.302	1.00	44.74	C
ATOM	310	OE1	GLU	A	54	30.546	30.379	4.509	1.00	46.67	O
ATOM	311	OE2	GLU	A	54	30.972	28.596	3.234	1.00	44.49	O
ATOM	312	N	LEU	A	55	29.812	28.468	9.424	1.00	38.37	N
ATOM	313	CA	LEU	A	55	29.291	29.137	10.583	1.00	38.08	C
ATOM	314	C	LEU	A	55	29.906	28.613	11.847	1.00	37.88	C
ATOM	315	O	LEU	A	55	30.359	29.397	12.693	1.00	37.59	O

ATOM	316	CB	LEU	A	55	27.789	29.006	10.648	1.00	38.28	
ATOM	317	CG	LEU	A	55	27.142	29.846	9.564	1.00	37.79	C
ATOM	318	CD1	LEU	A	55	25.722	29.443	9.409	1.00	38.31	C
ATOM	319	CD2	LEU	A	55	27.233	31.279	9.951	1.00	37.31	C
ATOM	320	N	ILE	A	56	29.938	27.286	11.971	1.00	37.71	C
ATOM	321	CA	ILE	A	56	30.455	26.667	13.183	1.00	37.35	N
ATOM	322	C	ILE	A	56	31.912	26.988	13.321	1.00	37.30	C
ATOM	323	O	ILE	A	56	32.373	27.320	14.406	1.00	36.89	C
ATOM	324	CB	ILE	A	56	30.288	25.171	13.175	1.00	37.30	O
ATOM	325	CG1	ILE	A	56	28.810	24.789	13.213	1.00	36.97	C
ATOM	326	CG2	ILE	A	56	30.968	24.602	14.397	1.00	37.75	C
ATOM	327	CD1	ILE	A	56	28.543	23.358	12.820	1.00	36.35	C
ATOM	328	N	GLU	A	57	32.627	26.899	12.204	1.00	37.67	C
ATOM	329	CA	GLU	A	57	34.045	27.204	12.166	1.00	38.21	N
ATOM	330	C	GLU	A	57	34.277	28.569	12.784	1.00	38.33	C
ATOM	331	O	GLU	A	57	35.210	28.758	13.566	1.00	38.33	C
ATOM	332	CB	GLU	A	57	34.547	27.214	10.725	1.00	38.49	O
ATOM	333	CG	GLU	A	57	36.038	27.461	10.559	1.00	40.09	C
ATOM	334	CD	GLU	A	57	36.870	26.539	11.424	1.00	42.96	C
ATOM	335	OE1	GLU	A	57	36.527	25.329	11.487	1.00	43.20	O
ATOM	336	OE2	GLU	A	57	37.852	27.032	12.044	1.00	44.14	O
ATOM	337	N	ASN	A	58	33.394	29.508	12.460	1.00	38.32	N
ATOM	338	CA	ASN	A	58	33.546	30.890	12.891	1.00	38.38	C
ATOM	339	C	ASN	A	58	32.849	31.221	14.167	1.00	38.06	C
ATOM	340	O	ASN	A	58	32.683	32.387	14.507	1.00	37.86	C
ATOM	341	CB	ASN	A	58	32.955	31.813	11.853	1.00	38.73	O
ATOM	342	CG	ASN	A	58	33.991	32.455	11.013	1.00	39.49	C
ATOM	343	OD1	ASN	A	58	34.501	31.841	10.083	1.00	42.51	C
ATOM	344	ND2	ASN	A	58	34.322	33.704	11.324	1.00	40.15	O
ATOM	345	N	GLU	A	59	32.399	30.196	14.856	1.00	37.88	N
ATOM	346	CA	GLU	A	59	31.683	30.396	16.092	1.00	37.68	C
ATOM	347	C	GLU	A	59	30.479	31.307	16.005	1.00	37.57	C
ATOM	348	O	GLU	A	59	30.305	32.203	16.813	1.00	36.94	O
ATOM	349	CB	GLU	A	59	32.663	30.857	17.118	1.00	37.60	C
ATOM	350	CG	GLU	A	59	33.710	29.785	17.212	1.00	38.25	C
ATOM	351	CD	GLU	A	59	34.545	29.886	18.435	1.00	38.10	C
ATOM	352	OE1	GLU	A	59	35.654	30.430	18.303	1.00	40.30	O
ATOM	353	OE2	GLU	A	59	34.089	29.420	19.498	1.00	37.62	O
ATOM	354	N	GLU	A	60	29.630	31.025	15.025	1.00	37.73	N
ATOM	355	CA	GLU	A	60	28.347	31.692	14.902	1.00	37.79	C
ATOM	356	C	GLU	A	60	27.290	30.620	14.923	1.00	37.12	C
ATOM	357	O	GLU	A	60	27.488	29.539	14.388	1.00	37.94	O
ATOM	358	CB	GLU	A	60	28.252	32.456	13.606	1.00	38.14	C
ATOM	359	CG	GLU	A	60	29.388	33.411	13.414	1.00	39.56	C
ATOM	360	CD	GLU	A	60	28.986	34.547	12.518	1.00	42.62	C
ATOM	361	OE1	GLU	A	60	29.023	34.389	11.276	1.00	42.22	O
ATOM	362	OE2	GLU	A	60	28.611	35.601	13.082	1.00	46.90	O
ATOM	363	N	PRO	A	61	26.158	30.917	15.517	1.00	36.25	N
ATOM	364	CA	PRO	A	61	25.111	29.919	15.680	1.00	36.11	C
ATOM	365	C	PRO	A	61	24.522	29.448	14.373	1.00	35.80	C
ATOM	366	O	PRO	A	61	24.534	30.159	13.386	1.00	36.29	O
ATOM	367	CB	PRO	A	61	24.021	30.672	16.442	1.00	36.35	C
ATOM	368	CG	PRO	A	61	24.367	32.117	16.353	1.00	36.05	C
ATOM	369	CD	PRO	A	61	25.790	32.228	16.061	1.00	36.10	C
ATOM	370	N	VAL	A	62	23.981	28.247	14.380	1.00	35.41	N
ATOM	371	CA	VAL	A	62	23.291	27.722	13.228	1.00	34.88	C
ATOM	372	C	VAL	A	62	22.438	26.584	13.720	1.00	34.68	C
ATOM	373	O	VAL	A	62	22.807	25.885	14.658	1.00	34.41	O
ATOM	374	CB	VAL	A	62	24.237	27.189	12.156	1.00	34.79	C
ATOM	375	CG1	VAL	A	62	25.108	26.062	12.695	1.00	34.66	C
ATOM	376	CG2	VAL	A	62	23.440	26.693	10.996	1.00	34.83	C

ATOM	377	N	VAL	A	63	21.271	26.418	13.118	1.00	34.54
ATOM	378	CA	VAL	A	63	20.454	25.273	13.447	1.00	34.18
ATOM	379	C	VAL	A	63	20.660	24.230	12.372	1.00	33.85
ATOM	380	O	VAL	A	63	20.486	24.512	11.203	1.00	33.36
ATOM	381	CB	VAL	A	63	18.980	25.622	13.523	1.00	34.05
ATOM	382	CG1	VAL	A	63	18.167	24.349	13.626	1.00	34.14
ATOM	383	CG2	VAL	A	63	18.717	26.502	14.710	1.00	33.51
ATOM	384	N	LEU	A	64	21.048	23.028	12.769	1.00	33.87
ATOM	385	CA	LEU	A	64	21.204	21.942	11.814	1.00	34.03
ATOM	386	C	LEU	A	64	19.894	21.193	11.856	1.00	33.70
ATOM	387	O	LEU	A	64	19.368	20.956	12.932	1.00	34.18
ATOM	388	CB	LEU	A	64	22.417	21.071	12.164	1.00	33.90
ATOM	389	CG	LEU	A	64	23.667	21.978	12.155	1.00	34.85
ATOM	390	CD1	LEU	A	64	24.975	21.303	12.485	1.00	35.82
ATOM	391	CD2	LEU	A	64	23.821	22.629	10.798	1.00	35.39
ATOM	392	N	THR	A	65	19.314	20.887	10.704	1.00	33.36
ATOM	393	CA	THR	A	65	18.035	20.177	10.704	1.00	33.24
ATOM	394	C	THR	A	65	18.105	18.683	10.517	1.00	32.87
ATOM	395	O	THR	A	65	17.096	18.016	10.715	1.00	32.11
ATOM	396	CB	THR	A	65	17.149	20.657	9.562	1.00	33.35
ATOM	397	OG1	THR	A	65	17.858	20.540	8.320	1.00	33.15
ATOM	398	CG2	THR	A	65	16.843	22.093	9.693	1.00	33.51
ATOM	399	N	ASP	A	66	19.265	18.173	10.106	1.00	33.09
ATOM	400	CA	ASP	A	66	19.392	16.767	9.724	1.00	33.53
ATOM	401	C	ASP	A	66	20.629	16.019	10.189	1.00	33.55
ATOM	402	O	ASP	A	66	21.136	15.179	9.458	1.00	33.68
ATOM	403	CB	ASP	A	66	19.339	16.653	8.192	1.00	33.55
ATOM	404	CG	ASP	A	66	20.397	17.484	7.508	1.00	34.00
ATOM	405	OD1	ASP	A	66	21.124	18.232	8.188	1.00	35.03
ATOM	406	OD2	ASP	A	66	20.583	17.456	6.284	1.00	35.31
ATOM	407	N	THR	A	67	21.107	16.264	11.397	1.00	33.83
ATOM	408	CA	THR	A	67	22.286	15.540	11.845	1.00	33.70
ATOM	409	C	THR	A	67	21.937	14.177	12.340	1.00	33.84
ATOM	410	O	THR	A	67	22.808	13.340	12.431	1.00	34.42
ATOM	411	CB	THR	A	67	22.951	16.215	13.020	1.00	33.66
ATOM	412	OG1	THR	A	67	21.981	16.451	14.046	1.00	33.90
ATOM	413	CG2	THR	A	67	23.472	17.562	12.674	1.00	33.56
ATOM	414	N	ASN	A	68	20.686	13.954	12.719	1.00	34.00
ATOM	415	CA	ASN	A	68	20.320	12.676	13.312	1.00	34.15
ATOM	416	C	ASN	A	68	21.108	12.454	14.575	1.00	33.97
ATOM	417	O	ASN	A	68	21.281	11.330	15.008	1.00	33.83
ATOM	418	CB	ASN	A	68	20.653	11.531	12.375	1.00	34.26
ATOM	419	CG	ASN	A	68	19.685	11.411	11.239	1.00	35.12
ATOM	420	OD1	ASN	A	68	18.512	11.104	11.434	1.00	35.89
ATOM	421	ND2	ASN	A	68	20.168	11.649	10.033	1.00	36.88
ATOM	422	N	LEU	A	69	21.598	13.531	15.162	1.00	34.12
ATOM	423	CA	LEU	A	69	22.446	13.419	16.334	1.00	34.19
ATOM	424	C	LEU	A	69	21.836	12.562	17.440	1.00	34.16
ATOM	425	O	LEU	A	69	22.538	11.737	18.039	1.00	33.75
ATOM	426	CB	LEU	A	69	22.774	14.808	16.881	1.00	34.13
ATOM	427	CG	LEU	A	69	23.652	14.795	18.125	1.00	34.10
ATOM	428	CD1	LEU	A	69	24.978	14.105	17.854	1.00	34.66
ATOM	429	CD2	LEU	A	69	23.883	16.183	18.582	1.00	33.94
ATOM	430	N	VAL	A	70	20.555	12.783	17.732	1.00	34.15
ATOM	431	CA	VAL	A	70	19.877	12.023	18.777	1.00	34.48
ATOM	432	C	VAL	A	70	18.654	11.307	18.228	1.00	34.64
ATOM	433	O	VAL	A	70	17.600	11.223	18.865	1.00	34.46
ATOM	434	CB	VAL	A	70	19.540	12.899	20.003	1.00	34.61
ATOM	435	CG1	VAL	A	70	20.808	13.555	20.519	1.00	35.04
ATOM	436	CG2	VAL	A	70	18.493	13.939	19.706	1.00	34.21
ATOM	437	N	TYR	A	71	18.833	10.762	17.033	1.00	34.98

N
C
C
O
C
C
C
N
C
C
O
C
C
C
N
C
C
O
C
C
O
C
O
N
C
C
O
C
C
C
N
C
C
O
C
C
C
N
C
C
O
C
C
C
N

ATOM	438	CA	TYR	A	71	17.779	10.041	16.352	1.00	35.19
ATOM	439	C	TYR	A	71	17.051	9.086	17.305	1.00	35.24
ATOM	440	O	TYR	A	71	15.837	9.148	17.421	1.00	35.61
ATOM	441	CB	TYR	A	71	18.337	9.291	15.137	1.00	35.26
ATOM	442	CG	TYR	A	71	17.352	8.278	14.618	1.00	35.95
ATOM	443	CD1	TYR	A	71	16.155	8.696	14.055	1.00	35.96
ATOM	444	CD2	TYR	A	71	17.584	6.915	14.731	1.00	34.66
ATOM	445	CE1	TYR	A	71	15.239	7.805	13.610	1.00	35.86
ATOM	446	CE2	TYR	A	71	16.664	6.011	14.281	1.00	34.47
ATOM	447	CZ	TYR	A	71	15.485	6.463	13.724	1.00	36.06
ATOM	448	OH	TYR	A	71	14.518	5.589	13.248	1.00	38.92
ATOM	449	N	PRO	A	72	17.774	8.211	17.993	1.00	35.15
ATOM	450	CA	PRO	A	72	17.141	7.271	18.924	1.00	35.16
ATOM	451	C	PRO	A	72	16.357	7.923	20.068	1.00	35.14
ATOM	452	O	PRO	A	72	15.473	7.294	20.643	1.00	34.97
ATOM	453	CB	PRO	A	72	18.327	6.475	19.487	1.00	35.18
ATOM	454	CG	PRO	A	72	19.433	6.677	18.524	1.00	35.26
ATOM	455	CD	PRO	A	72	19.233	8.031	17.938	1.00	35.29
ATOM	456	N	ALA	A	73	16.657	9.169	20.401	1.00	35.18
ATOM	457	CA	ALA	A	73	15.967	9.803	21.512	1.00	35.21
ATOM	458	C	ALA	A	73	14.657	10.414	21.097	1.00	35.08
ATOM	459	O	ALA	A	73	13.890	10.862	21.934	1.00	34.86
ATOM	460	CB	ALA	A	73	16.829	10.864	22.135	1.00	35.37
ATOM	461	N	LEU	A	74	14.370	10.434	19.812	1.00	35.25
ATOM	462	CA	LEU	A	74	13.142	11.090	19.399	1.00	35.54
ATOM	463	C	LEU	A	74	11.903	10.434	19.958	1.00	35.66
ATOM	464	O	LEU	A	74	10.893	11.088	20.118	1.00	35.64
ATOM	465	CB	LEU	A	74	13.044	11.190	17.894	1.00	35.31
ATOM	466	CG	LEU	A	74	14.161	12.046	17.333	1.00	36.48
ATOM	467	CD1	LEU	A	74	13.887	12.277	15.890	1.00	37.76
ATOM	468	CD2	LEU	A	74	14.300	13.394	18.076	1.00	37.06
ATOM	469	N	LYS	A	75	11.979	9.151	20.282	1.00	36.19
ATOM	470	CA	LYS	A	75	10.795	8.453	20.767	1.00	36.36
ATOM	471	C	LYS	A	75	10.566	8.710	22.246	1.00	36.85
ATOM	472	O	LYS	A	75	9.491	8.437	22.765	1.00	37.12
ATOM	473	CB	LYS	A	75	10.904	6.956	20.505	1.00	36.04
ATOM	474	CG	LYS	A	75	12.060	6.284	21.216	1.00	35.77
ATOM	475	CD	LYS	A	75	12.245	4.829	20.782	1.00	34.46
ATOM	476	CE	LYS	A	75	13.720	4.460	20.710	1.00	33.42
ATOM	477	NZ	LYS	A	75	14.375	4.536	22.036	1.00	32.34
ATOM	478	N	TRP	A	76	11.578	9.244	22.920	1.00	37.24
ATOM	479	CA	TRP	A	76	11.481	9.510	24.345	1.00	37.30
ATOM	480	C	TRP	A	76	10.274	10.369	24.700	1.00	37.42
ATOM	481	O	TRP	A	76	9.917	11.306	23.984	1.00	37.54
ATOM	482	CB	TRP	A	76	12.728	10.247	24.838	1.00	37.12
ATOM	483	CG	TRP	A	76	13.980	9.446	24.816	1.00	36.77
ATOM	484	CD1	TRP	A	76	14.142	8.176	24.359	1.00	37.03
ATOM	485	CD2	TRP	A	76	15.255	9.864	25.291	1.00	36.31
ATOM	486	NE1	TRP	A	76	15.445	7.776	24.527	1.00	36.52
ATOM	487	CE2	TRP	A	76	16.147	8.801	25.097	1.00	36.08
ATOM	488	CE3	TRP	A	76	15.737	11.040	25.864	1.00	36.23
ATOM	489	CZ2	TRP	A	76	17.480	8.881	25.443	1.00	36.27
ATOM	490	CZ3	TRP	A	76	17.055	11.113	26.209	1.00	36.35
ATOM	491	CH2	TRP	A	76	17.915	10.045	25.996	1.00	36.41
ATOM	492	N	ASP	A	77	9.662	10.015	25.820	1.00	37.42
ATOM	493	CA	ASP	A	77	8.591	10.768	26.435	1.00	37.36
ATOM	494	C	ASP	A	77	8.707	10.370	27.907	1.00	37.09
ATOM	495	O	ASP	A	77	9.628	9.647	28.260	1.00	37.04
ATOM	496	CB	ASP	A	77	7.245	10.377	25.834	1.00	37.48
ATOM	497	CG	ASP	A	77	6.990	8.888	25.903	1.00	38.01
ATOM	498	OD1	ASP	A	77	7.704	8.191	26.658	1.00	38.42

[illegible]

ATOM	499	OD2	ASP	A	77	6.093	8.319	25.244	1.00	39.67	O
ATOM	500	N	LEU	A	78	7.789	10.798	28.763	1.00	36.79	N
ATOM	501	CA	LEU	A	78	7.913	10.496	30.182	1.00	36.49	C
ATOM	502	C	LEU	A	78	7.729	9.026	30.464	1.00	36.62	C
ATOM	503	O	LEU	A	78	8.456	8.426	31.260	1.00	36.50	O
ATOM	504	CB	LEU	A	78	6.903	11.296	30.974	1.00	36.33	C
ATOM	505	CG	LEU	A	78	7.119	12.796	30.863	1.00	36.49	C
ATOM	506	CD1	LEU	A	78	6.008	13.510	31.594	1.00	36.63	C
ATOM	507	CD2	LEU	A	78	8.494	13.194	31.399	1.00	36.10	C
ATOM	508	N	GLU	A	79	6.748	8.433	29.815	1.00	36.81	N
ATOM	509	CA	GLU	A	79	6.505	7.030	30.041	1.00	36.97	C
ATOM	510	C	GLU	A	79	7.737	6.176	29.738	1.00	36.75	C
ATOM	511	O	GLU	A	79	8.151	5.373	30.572	1.00	36.84	O
ATOM	512	CB	GLU	A	79	5.323	6.560	29.214	1.00	37.25	C
ATOM	513	CG	GLU	A	79	4.937	5.137	29.540	1.00	37.93	C
ATOM	514	CD	GLU	A	79	3.729	4.681	28.768	1.00	38.61	C
ATOM	515	OE1	GLU	A	79	3.381	5.352	27.775	1.00	38.32	O
ATOM	516	OE2	GLU	A	79	3.131	3.654	29.166	1.00	40.10	O
ATOM	517	N	TYR	A	80	8.318	6.337	28.553	1.00	36.50	N
ATOM	518	CA	TYR	A	80	9.489	5.549	28.176	1.00	36.32	C
ATOM	519	C	TYR	A	80	10.644	5.804	29.123	1.00	36.27	C
ATOM	520	O	TYR	A	80	11.343	4.872	29.516	1.00	36.15	O
ATOM	521	CB	TYR	A	80	9.921	5.889	26.751	1.00	36.39	C
ATOM	522	CG	TYR	A	80	11.180	5.202	26.228	1.00	36.06	C
ATOM	523	CD1	TYR	A	80	11.100	4.000	25.557	1.00	36.22	C
ATOM	524	CD2	TYR	A	80	12.432	5.793	26.349	1.00	35.91	C
ATOM	525	CE1	TYR	A	80	12.225	3.380	25.051	1.00	36.52	C
ATOM	526	CE2	TYR	A	80	13.568	5.179	25.844	1.00	36.04	C
ATOM	527	CZ	TYR	A	80	13.454	3.967	25.190	1.00	36.46	C
ATOM	528	OH	TYR	A	80	14.561	3.321	24.673	1.00	35.81	O
ATOM	529	N	LEU	A	81	10.846	7.064	29.493	1.00	36.13	N
ATOM	530	CA	LEU	A	81	11.971	7.408	30.353	1.00	36.22	C
ATOM	531	C	LEU	A	81	11.777	6.827	31.747	1.00	36.13	C
ATOM	532	O	LEU	A	81	12.706	6.259	32.325	1.00	35.90	O
ATOM	533	CB	LEU	A	81	12.208	8.930	30.406	1.00	36.22	C
ATOM	534	CG	LEU	A	81	12.774	9.568	29.121	1.00	36.35	C
ATOM	535	CD1	LEU	A	81	12.880	11.055	29.274	1.00	36.49	C
ATOM	536	CD2	LEU	A	81	14.132	9.017	28.718	1.00	36.18	C
ATOM	537	N	GLN	A	82	10.569	6.956	32.280	1.00	36.15	N
ATOM	538	CA	GLN	A	82	10.284	6.424	33.597	1.00	36.29	C
ATOM	539	C	GLN	A	82	10.575	4.927	33.605	1.00	36.25	C
ATOM	540	O	GLN	A	82	11.210	4.408	34.515	1.00	35.96	O
ATOM	541	CB	GLN	A	82	8.838	6.710	33.976	1.00	36.38	C
ATOM	542	CG	GLN	A	82	8.418	6.080	35.279	1.00	37.00	C
ATOM	543	CD	GLN	A	82	7.191	6.740	35.872	1.00	37.85	C
ATOM	544	OE1	GLN	A	82	6.640	7.676	35.295	1.00	38.24	O
ATOM	545	NE2	GLN	A	82	6.754	6.247	37.020	1.00	38.93	N
ATOM	546	N	GLU	A	83	10.136	4.240	32.560	1.00	36.46	N
ATOM	547	CA	GLU	A	83	10.366	2.807	32.451	1.00	36.59	C
ATOM	548	C	GLU	A	83	11.843	2.424	32.307	1.00	36.50	C
ATOM	549	O	GLU	A	83	12.228	1.321	32.677	1.00	36.55	O
ATOM	550	CB	GLU	A	83	9.568	2.237	31.271	1.00	36.57	C
ATOM	551	CG	GLU	A	83	9.497	0.714	31.239	1.00	36.95	C
ATOM	552	CD	GLU	A	83	8.732	0.107	32.416	1.00	37.61	C
ATOM	553	OE1	GLU	A	83	7.922	0.815	33.050	1.00	37.88	O
ATOM	554	OE2	GLU	A	83	8.938	-1.090	32.719	1.00	37.54	O
ATOM	555	N	ASN	A	84	12.680	3.324	31.805	1.00	36.58	N
ATOM	556	CA	ASN	A	84	14.049	2.936	31.476	1.00	36.62	C
ATOM	557	C	ASN	A	84	15.210	3.749	32.001	1.00	36.66	C
ATOM	558	O	ASN	A	84	16.349	3.310	31.900	1.00	36.62	O
ATOM	559	CB	ASN	A	84	14.211	2.936	29.960	1.00	36.62	C

ATOM	560	CG	ASN	A	84	13.362	1.902	29.286	1.00	36.37
ATOM	561	OD1	ASN	A	84	13.352	0.733	29.683	1.00	36.46
ATOM	562	ND2	ASN	A	84	12.644	2.317	28.247	1.00	35.65
ATOM	563	N	ILE	A	85	14.975	4.920	32.557	1.00	36.80
ATOM	564	CA	ILE	A	85	16.125	5.730	32.885	1.00	37.13
ATOM	565	C	ILE	A	85	16.834	5.365	34.180	1.00	37.28
ATOM	566	O	ILE	A	85	17.793	6.022	34.568	1.00	37.74
ATOM	567	CB	ILE	A	85	15.752	7.184	32.874	1.00	37.21
ATOM	568	CG1	ILE	A	85	16.958	8.009	32.445	1.00	37.55
ATOM	569	CG2	ILE	A	85	15.241	7.593	34.222	1.00	37.38
ATOM	570	CD1	ILE	A	85	16.623	9.467	32.195	1.00	37.68
ATOM	571	N	GLY	A	86	16.383	4.326	34.855	1.00	37.29
ATOM	572	CA	GLY	A	86	17.089	3.892	36.038	1.00	37.39
ATOM	573	C	GLY	A	86	16.553	4.496	37.308	1.00	37.43
ATOM	574	O	GLY	A	86	15.583	5.265	37.290	1.00	37.34
ATOM	575	N	ASN	A	87	17.212	4.167	38.414	1.00	37.34
ATOM	576	CA	ASN	A	87	16.750	4.602	39.716	1.00	37.42
ATOM	577	C	ASN	A	87	17.701	5.603	40.357	1.00	37.39
ATOM	578	O	ASN	A	87	17.740	5.742	41.578	1.00	37.39
ATOM	579	CB	ASN	A	87	16.545	3.399	40.612	1.00	37.40
ATOM	580	N	GLY	A	88	18.451	6.321	39.529	1.00	37.47
ATOM	581	CA	GLY	A	88	19.405	7.296	40.028	1.00	37.41
ATOM	582	C	GLY	A	88	18.686	8.554	40.450	1.00	37.30
ATOM	583	O	GLY	A	88	17.500	8.709	40.171	1.00	37.36
ATOM	584	N	ASP	A	89	19.390	9.453	41.125	1.00	37.26
ATOM	585	CA	ASP	A	89	18.780	10.716	41.531	1.00	37.21
ATOM	586	C	ASP	A	89	18.726	11.681	40.355	1.00	36.91
ATOM	587	O	ASP	A	89	19.607	11.654	39.500	1.00	36.62
ATOM	588	CB	ASP	A	89	19.563	11.346	42.680	1.00	37.24
ATOM	589	CG	ASP	A	89	19.277	10.690	44.000	1.00	37.24
ATOM	590	OD1	ASP	A	89	18.398	9.802	44.044	1.00	37.14
ATOM	591	OD2	ASP	A	89	19.876	11.002	45.047	1.00	37.98
ATOM	592	N	PHE	A	90	17.667	12.492	40.295	1.00	36.88
ATOM	593	CA	PHE	A	90	17.554	13.570	39.298	1.00	36.87
ATOM	594	C	PHE	A	90	17.327	14.926	39.942	1.00	36.79
ATOM	595	O	PHE	A	90	16.455	15.080	40.795	1.00	36.69
ATOM	596	CB	PHE	A	90	16.420	13.308	38.321	1.00	36.70
ATOM	597	CG	PHE	A	90	16.712	12.210	37.371	1.00	36.83
ATOM	598	CD1	PHE	A	90	16.551	10.901	37.757	1.00	36.07
ATOM	599	CD2	PHE	A	90	17.191	12.479	36.102	1.00	37.27
ATOM	600	CE1	PHE	A	90	16.827	9.885	36.899	1.00	35.81
ATOM	601	CE2	PHE	A	90	17.475	11.449	35.238	1.00	36.82
ATOM	602	CZ	PHE	A	90	17.291	10.150	35.642	1.00	36.06
ATOM	603	N	SER	A	91	18.123	15.905	39.529	1.00	36.92
ATOM	604	CA	SER	A	91	17.971	17.266	40.027	1.00	36.99
ATOM	605	C	SER	A	91	16.738	17.923	39.428	1.00	37.10
ATOM	606	O	SER	A	91	16.586	17.985	38.206	1.00	36.78
ATOM	607	CB	SER	A	91	19.200	18.105	39.699	1.00	36.86
ATOM	608	OG	SER	A	91	20.350	17.616	40.358	1.00	36.54
ATOM	609	N	VAL	A	92	15.857	18.402	40.303	1.00	37.42
ATOM	610	CA	VAL	A	92	14.660	19.106	39.876	1.00	37.75
ATOM	611	C	VAL	A	92	14.509	20.468	40.545	1.00	38.10
ATOM	612	O	VAL	A	92	14.472	20.589	41.768	1.00	38.08
ATOM	613	CB	VAL	A	92	13.406	18.303	40.159	1.00	37.63
ATOM	614	CG1	VAL	A	92	12.197	19.034	39.621	1.00	37.42
ATOM	615	CG2	VAL	A	92	13.515	16.946	39.527	1.00	37.78
ATOM	616	N	TYR	A	93	14.384	21.487	39.709	1.00	38.59
ATOM	617	CA	TYR	A	93	14.220	22.849	40.166	1.00	38.91
ATOM	618	C	TYR	A	93	12.784	23.173	40.219	1.00	39.41
ATOM	619	O	TYR	A	93	12.019	22.786	39.343	1.00	39.64
ATOM	620	CB	TYR	A	93	14.883	23.799	39.202	1.00	38.84

C
O
N
N
C
C
O
C
C
C
C
N
C
C
O
N
C
C
C
O
N
C
C
C
O
C
C
C
C
N
C
C
C
O
C
O
C

ATOM	621	CG	TYR	A	93	16.332	23.588	39.288	1.00	38.25	C
ATOM	622	CD1	TYR	A	93	17.044	24.122	40.334	1.00	38.21	C
ATOM	623	CD2	TYR	A	93	16.968	22.750	38.415	1.00	37.50	C
ATOM	624	CE1	TYR	A	93	18.362	23.893	40.464	1.00	37.90	C
ATOM	625	CE2	TYR	A	93	18.287	22.512	38.534	1.00	37.82	C
ATOM	626	CZ	TYR	A	93	18.987	23.090	39.557	1.00	37.78	C
ATOM	627	OH	TYR	A	93	20.322	22.836	39.677	1.00	39.39	O
ATOM	628	N	SER	A	94	12.422	23.935	41.228	1.00	40.00	N
ATOM	629	CA	SER	A	94	11.043	24.274	41.416	1.00	40.51	C
ATOM	630	C	SER	A	94	11.020	25.773	41.426	1.00	40.68	C
ATOM	631	O	SER	A	94	11.962	26.384	41.896	1.00	41.00	O
ATOM	632	CB	SER	A	94	10.539	23.693	42.730	1.00	40.58	C
ATOM	633	OG	SER	A	94	9.139	23.874	42.858	1.00	41.30	O
ATOM	634	N	ALA	A	95	9.974	26.374	40.878	1.00	40.87	N
ATOM	635	CA	ALA	A	95	9.899	27.824	40.856	1.00	40.91	C
ATOM	636	C	ALA	A	95	8.483	28.343	40.826	1.00	41.03	C
ATOM	637	O	ALA	A	95	7.573	27.699	40.309	1.00	41.09	O
ATOM	638	CB	ALA	A	95	10.630	28.350	39.668	1.00	40.93	C
ATOM	639	N	SER	A	96	8.328	29.549	41.350	1.00	41.12	N
ATOM	640	CA	SER	A	96	7.034	30.200	41.413	1.00	41.05	C
ATOM	641	C	SER	A	96	6.834	31.208	40.305	1.00	40.72	C
ATOM	642	O	SER	A	96	5.833	31.912	40.277	1.00	40.86	O
ATOM	643	CB	SER	A	96	6.902	30.938	42.731	1.00	41.16	C
ATOM	644	OG	SER	A	96	5.767	31.779	42.693	1.00	41.90	O
ATOM	645	N	THR	A	97	7.808	31.314	39.423	1.00	40.34	N
ATOM	646	CA	THR	A	97	7.710	32.209	38.292	1.00	40.18	C
ATOM	647	C	THR	A	97	8.073	31.368	37.113	1.00	39.72	C
ATOM	648	O	THR	A	97	8.582	30.274	37.279	1.00	39.83	O
ATOM	649	CB	THR	A	97	8.732	33.336	38.393	1.00	40.49	C
ATOM	650	OG1	THR	A	97	8.905	33.946	37.105	1.00	41.00	O
ATOM	651	CG2	THR	A	97	10.145	32.779	38.718	1.00	40.86	C
ATOM	652	N	HIS	A	98	7.862	31.874	35.915	1.00	39.32	N
ATOM	653	CA	HIS	A	98	8.257	31.112	34.754	1.00	39.12	C
ATOM	654	C	HIS	A	98	9.765	31.163	34.543	1.00	39.41	C
ATOM	655	O	HIS	A	98	10.299	30.423	33.711	1.00	39.15	O
ATOM	656	CB	HIS	A	98	7.576	31.646	33.516	1.00	38.95	C
ATOM	657	CG	HIS	A	98	7.807	33.104	33.290	1.00	38.64	C
ATOM	658	ND1	HIS	A	98	7.095	34.081	33.950	1.00	37.36	N
ATOM	659	CD2	HIS	A	98	8.672	33.753	32.477	1.00	38.57	C
ATOM	660	CE1	HIS	A	98	7.509	35.268	33.550	1.00	37.33	C
ATOM	661	NE2	HIS	A	98	8.463	35.098	32.654	1.00	37.37	N
ATOM	662	N	LYS	A	99	10.452	32.021	35.294	1.00	39.52	N
ATOM	663	CA	LYS	A	99	11.881	32.184	35.105	1.00	39.98	C
ATOM	664	C	LYS	A	99	12.749	31.364	36.029	1.00	40.38	C
ATOM	665	O	LYS	A	99	12.744	31.554	37.246	1.00	40.33	O
ATOM	666	CB	LYS	A	99	12.281	33.631	35.297	1.00	40.10	C
ATOM	667	CG	LYS	A	99	11.814	34.527	34.219	1.00	40.15	C
ATOM	668	CD	LYS	A	99	12.537	35.820	34.337	1.00	40.36	C
ATOM	669	CE	LYS	A	99	11.725	36.835	35.072	1.00	41.10	C
ATOM	670	NZ	LYS	A	99	10.975	37.655	34.074	1.00	41.52	N
ATOM	671	N	PHE	A	100	13.541	30.486	35.433	1.00	40.77	N
ATOM	672	CA	PHE	A	100	14.450	29.665	36.201	1.00	41.12	C
ATOM	673	C	PHE	A	100	15.893	30.159	36.218	1.00	41.78	C
ATOM	674	O	PHE	A	100	16.808	29.463	35.765	1.00	42.13	O
ATOM	675	CB	PHE	A	100	14.418	28.255	35.658	1.00	40.86	C
ATOM	676	CG	PHE	A	100	13.211	27.494	36.060	1.00	40.53	C
ATOM	677	CD1	PHE	A	100	12.040	27.604	35.344	1.00	39.76	C
ATOM	678	CD2	PHE	A	100	13.249	26.666	37.159	1.00	39.54	C
ATOM	679	CE1	PHE	A	100	10.942	26.890	35.707	1.00	39.56	C
ATOM	680	CE2	PHE	A	100	12.155	25.955	37.526	1.00	39.79	C
ATOM	681	CZ	PHE	A	100	10.995	26.064	36.802	1.00	39.91	C

ATOM	682	N	LEU	A	101	16.124	31.347	36.747	1.00	42.23	N
ATOM	683	CA	LEU	A	101	17.501	31.800	36.868	1.00	42.74	C
ATOM	684	C	LEU	A	101	18.429	30.780	37.567	1.00	43.27	C
ATOM	685	O	LEU	A	101	18.213	30.392	38.736	1.00	43.22	O
ATOM	686	CB	LEU	A	101	17.549	33.097	37.657	1.00	42.90	C
ATOM	687	CG	LEU	A	101	18.900	33.800	37.793	1.00	43.35	C
ATOM	688	CD1	LEU	A	101	19.441	34.288	36.455	1.00	43.59	C
ATOM	689	CD2	LEU	A	101	18.703	34.967	38.724	1.00	43.91	C
ATOM	690	N	TYR	A	102	19.470	30.379	36.836	1.00	43.59	N
ATOM	691	CA	TYR	A	102	20.547	29.522	37.340	1.00	43.71	C
ATOM	692	C	TYR	A	102	21.380	30.186	38.460	1.00	43.57	C
ATOM	693	O	TYR	A	102	21.746	31.364	38.343	1.00	44.09	O
ATOM	694	CB	TYR	A	102	21.531	29.218	36.193	1.00	43.90	C
ATOM	695	CG	TYR	A	102	22.703	28.421	36.700	1.00	44.68	C
ATOM	696	CD1	TYR	A	102	22.553	27.080	37.033	1.00	44.29	C
ATOM	697	CD2	TYR	A	102	23.935	29.023	36.918	1.00	44.88	C
ATOM	698	CE1	TYR	A	102	23.602	26.358	37.541	1.00	44.98	C
ATOM	699	CE2	TYR	A	102	24.987	28.309	37.422	1.00	45.44	C
ATOM	700	CZ	TYR	A	102	24.826	26.984	37.735	1.00	45.86	C
ATOM	701	OH	TYR	A	102	25.922	26.313	38.237	1.00	47.07	O
ATOM	702	N	TYR	A	103	21.704	29.451	39.523	1.00	42.91	N
ATOM	703	CA	TYR	A	103	22.642	29.970	40.527	1.00	42.67	C
ATOM	704	C	TYR	A	103	23.561	28.895	41.102	1.00	41.98	C
ATOM	705	O	TYR	A	103	23.165	27.768	41.351	1.00	41.99	O
ATOM	706	CB	TYR	A	103	21.922	30.638	41.670	1.00	42.83	C
ATOM	707	CG	TYR	A	103	20.974	29.692	42.265	1.00	44.33	C
ATOM	708	CD1	TYR	A	103	19.770	29.454	41.646	1.00	46.69	C
ATOM	709	CD2	TYR	A	103	21.306	28.959	43.390	1.00	45.99	C
ATOM	710	CE1	TYR	A	103	18.880	28.539	42.147	1.00	47.83	C
ATOM	711	CE2	TYR	A	103	20.426	28.045	43.917	1.00	47.23	C
ATOM	712	CZ	TYR	A	103	19.203	27.836	43.284	1.00	48.65	C
ATOM	713	OH	TYR	A	103	18.287	26.921	43.774	1.00	51.36	O
ATOM	714	N	ASP	A	104	24.786	29.300	41.369	1.00	41.25	N
ATOM	715	CA	ASP	A	104	25.829	28.399	41.764	1.00	40.65	C
ATOM	716	C	ASP	A	104	25.924	28.335	43.267	1.00	40.45	C
ATOM	717	O	ASP	A	104	26.430	29.240	43.924	1.00	40.13	O
ATOM	718	CB	ASP	A	104	27.110	28.904	41.142	1.00	40.60	C
ATOM	719	CG	ASP	A	104	28.306	28.117	41.548	1.00	41.02	C
ATOM	720	OD1	ASP	A	104	28.270	27.482	42.629	1.00	42.21	O
ATOM	721	OD2	ASP	A	104	29.337	28.095	40.838	1.00	40.68	O
ATOM	722	N	GLU	A	105	25.446	27.222	43.801	1.00	40.41	N
ATOM	723	CA	GLU	A	105	25.368	27.029	45.231	1.00	40.21	C
ATOM	724	C	GLU	A	105	26.734	27.234	45.895	1.00	39.89	C
ATOM	725	O	GLU	A	105	26.803	27.738	47.021	1.00	39.65	O
ATOM	726	CB	GLU	A	105	24.769	25.643	45.528	1.00	40.44	C
ATOM	727	CG	GLU	A	105	23.247	25.572	45.351	1.00	41.43	C
ATOM	728	CD	GLU	A	105	22.708	24.150	45.207	1.00	42.48	C
ATOM	729	OE1	GLU	A	105	23.138	23.269	45.986	1.00	43.10	O
ATOM	730	OE2	GLU	A	105	21.842	23.914	44.322	1.00	42.00	O
ATOM	731	N	LYS	A	106	27.824	26.899	45.204	1.00	39.58	N
ATOM	732	CA	LYS	A	106	29.141	27.042	45.831	1.00	39.58	C
ATOM	733	C	LYS	A	106	29.427	28.469	46.190	1.00	39.64	C
ATOM	734	O	LYS	A	106	30.165	28.730	47.128	1.00	39.93	O
ATOM	735	CB	LYS	A	106	30.302	26.592	44.945	1.00	39.49	C
ATOM	736	CG	LYS	A	106	30.444	25.096	44.731	1.00	39.29	C
ATOM	737	N	LYS	A	107	28.856	29.397	45.438	1.00	39.68	N
ATOM	738	CA	LYS	A	107	29.140	30.794	45.663	1.00	39.58	C
ATOM	739	C	LYS	A	107	28.151	31.452	46.607	1.00	40.18	C
ATOM	740	O	LYS	A	107	28.279	32.637	46.883	1.00	40.30	O
ATOM	741	CB	LYS	A	107	29.161	31.531	44.325	1.00	39.46	C
ATOM	742	CG	LYS	A	107	30.390	31.199	43.455	1.00	38.46	C

ATOM	743	CD	LYS	A	107	30.299	31.737	42.029	1.00	36.98	C
ATOM	744	CE	LYS	A	107	31.643	31.632	41.292	1.00	36.36	C
ATOM	745	NZ	LYS	A	107	31.553	31.935	39.819	1.00	35.12	N
ATOM	746	N	MET	A	108	27.197	30.697	47.148	1.00	40.92	N
ATOM	747	CA	MET	A	108	26.165	31.316	47.978	1.00	41.42	C
ATOM	748	C	MET	A	108	26.675	31.937	49.258	1.00	42.03	C
ATOM	749	O	MET	A	108	26.142	32.952	49.704	1.00	42.33	O
ATOM	750	CB	MET	A	108	25.006	30.366	48.210	1.00	41.31	C
ATOM	751	CG	MET	A	108	24.285	30.179	46.892	1.00	41.95	C
ATOM	752	SD	MET	A	108	22.861	29.100	46.846	1.00	43.29	S
ATOM	753	CE	MET	A	108	21.737	30.010	48.064	1.00	43.04	C
ATOM	754	N	ALA	A	109	27.737	31.382	49.816	1.00	42.84	N
ATOM	755	CA	ALA	A	109	28.282	31.912	51.064	1.00	43.65	C
ATOM	756	C	ALA	A	109	28.655	33.387	50.963	1.00	44.40	C
ATOM	757	O	ALA	A	109	28.321	34.195	51.823	1.00	44.44	O
ATOM	758	CB	ALA	A	109	29.486	31.107	51.476	1.00	43.66	C
ATOM	759	N	ASN	A	110	29.332	33.729	49.882	1.00	45.46	N
ATOM	760	CA	ASN	A	110	29.817	35.084	49.665	1.00	46.25	C
ATOM	761	C	ASN	A	110	28.789	36.154	49.317	1.00	46.26	C
ATOM	762	O	ASN	A	110	29.163	37.304	49.066	1.00	46.51	O
ATOM	763	CB	ASN	A	110	30.841	35.036	48.543	1.00	46.62	C
ATOM	764	CG	ASN	A	110	32.024	34.145	48.886	1.00	47.94	C
ATOM	765	OD1	ASN	A	110	32.474	34.093	50.045	1.00	48.12	O
ATOM	766	ND2	ASN	A	110	32.530	33.432	47.883	1.00	48.75	N
ATOM	767	N	PHE	A	111	27.517	35.786	49.228	1.00	46.12	N
ATOM	768	CA	PHE	A	111	26.470	36.795	49.057	1.00	45.97	C
ATOM	769	C	PHE	A	111	25.267	36.385	49.889	1.00	46.17	C
ATOM	770	O	PHE	A	111	24.217	36.002	49.361	1.00	46.25	O
ATOM	771	CB	PHE	A	111	26.072	36.978	47.599	1.00	45.55	C
ATOM	772	CG	PHE	A	111	27.076	37.718	46.777	1.00	44.32	C
ATOM	773	CD1	PHE	A	111	28.269	37.123	46.419	1.00	43.65	C
ATOM	774	CD2	PHE	A	111	26.808	39.000	46.320	1.00	44.04	C
ATOM	775	CE1	PHE	A	111	29.187	37.794	45.620	1.00	43.19	C
ATOM	776	CE2	PHE	A	111	27.715	39.673	45.522	1.00	43.22	C
ATOM	777	CZ	PHE	A	111	28.911	39.068	45.177	1.00	43.00	C
ATOM	778	N	GLN	A	112	25.434	36.479	51.200	1.00	46.35	N
ATOM	779	CA	GLN	A	112	24.401	36.052	52.130	1.00	46.52	C
ATOM	780	C	GLN	A	112	23.040	36.638	51.754	1.00	46.68	C
ATOM	781	O	GLN	A	112	22.019	35.995	51.968	1.00	46.87	O
ATOM	782	CB	GLN	A	112	24.774	36.438	53.567	1.00	46.62	C
ATOM	783	CG	GLN	A	112	26.201	36.060	53.998	1.00	46.96	C
ATOM	784	CD	GLN	A	112	26.418	34.554	54.125	1.00	48.05	C
ATOM	785	OE1	GLN	A	112	25.834	33.762	53.372	1.00	47.89	O
ATOM	786	NE2	GLN	A	112	27.271	34.158	55.075	1.00	48.55	N
ATOM	787	N	ASN	A	113	23.019	37.836	51.168	1.00	46.80	N
ATOM	788	CA	ASN	A	113	21.750	38.463	50.793	1.00	46.89	C
ATOM	789	C	ASN	A	113	21.104	37.951	49.517	1.00	47.08	C
ATOM	790	O	ASN	A	113	20.105	38.521	49.077	1.00	47.55	O
ATOM	791	CB	ASN	A	113	21.898	39.976	50.682	1.00	46.69	C
ATOM	792	CG	ASN	A	113	22.258	40.604	52.001	1.00	46.91	C
ATOM	793	OD1	ASN	A	113	22.085	39.982	53.055	1.00	46.08	O
ATOM	794	ND2	ASN	A	113	22.767	41.837	51.962	1.00	46.59	N
ATOM	795	N	PHE	A	114	21.651	36.908	48.903	1.00	47.08	N
ATOM	796	CA	PHE	A	114	21.012	36.379	47.706	1.00	47.01	C
ATOM	797	C	PHE	A	114	19.986	35.364	48.145	1.00	47.11	C
ATOM	798	O	PHE	A	114	20.293	34.449	48.916	1.00	47.34	O
ATOM	799	CB	PHE	A	114	21.995	35.713	46.764	1.00	46.83	C
ATOM	800	CG	PHE	A	114	21.342	35.132	45.555	1.00	47.03	C
ATOM	801	CD1	PHE	A	114	20.674	35.953	44.655	1.00	47.22	C
ATOM	802	CD2	PHE	A	114	21.376	33.761	45.318	1.00	47.97	C
ATOM	803	CE1	PHE	A	114	20.060	35.422	43.522	1.00	47.53	C

ATOM	804	CE2	PHE	A	114	20.763	33.212	44.177	1.00	47.83	C
ATOM	805	CZ	PHE	A	114	20.104	34.044	43.282	1.00	47.67	C
ATOM	806	N	LYS	A	115	18.763	35.530	47.662	1.00	46.94	N
ATOM	807	CA	LYS	A	115	17.683	34.628	48.029	1.00	46.64	C
ATOM	808	C	LYS	A	115	17.109	34.063	46.744	1.00	46.38	C
ATOM	809	O	LYS	A	115	16.351	34.744	46.046	1.00	46.75	O
ATOM	810	CB	LYS	A	115	16.613	35.372	48.835	1.00	46.60	C
ATOM	811	N	PRO	A	116	17.445	32.813	46.449	1.00	45.84	N
ATOM	812	CA	PRO	A	116	17.079	32.192	45.165	1.00	45.70	C
ATOM	813	C	PRO	A	116	15.576	32.139	44.937	1.00	45.44	C
ATOM	814	O	PRO	A	116	14.827	31.956	45.892	1.00	45.46	O
ATOM	815	CB	PRO	A	116	17.640	30.761	45.260	1.00	45.71	C
ATOM	816	CG	PRO	A	116	18.435	30.689	46.536	1.00	45.66	C
ATOM	817	CD	PRO	A	116	18.137	31.890	47.359	1.00	45.75	C
ATOM	818	N	ARG	A	117	15.154	32.307	43.688	1.00	45.27	N
ATOM	819	CA	ARG	A	117	13.742	32.232	43.332	1.00	45.16	C
ATOM	820	C	ARG	A	117	13.291	30.791	43.104	1.00	45.12	C
ATOM	821	O	ARG	A	117	12.096	30.500	43.207	1.00	45.48	O
ATOM	822	CB	ARG	A	117	13.467	33.059	42.097	1.00	45.26	C
ATOM	823	N	SER	A	118	14.238	29.902	42.796	1.00	44.78	N
ATOM	824	CA	SER	A	118	13.944	28.480	42.594	1.00	44.60	C
ATOM	825	C	SER	A	118	14.782	27.583	43.509	1.00	44.63	C
ATOM	826	O	SER	A	118	15.925	27.921	43.816	1.00	45.32	O
ATOM	827	CB	SER	A	118	14.303	28.072	41.184	1.00	44.44	C
ATOM	828	OG	SER	A	118	15.651	27.624	41.186	1.00	44.35	O
ATOM	829	N	ASN	A	119	14.241	26.428	43.899	1.00	44.11	N
ATOM	830	CA	ASN	A	119	14.969	25.472	44.728	1.00	43.72	C
ATOM	831	C	ASN	A	119	15.295	24.196	43.953	1.00	43.16	C
ATOM	832	O	ASN	A	119	14.576	23.821	43.030	1.00	43.23	O
ATOM	833	CB	ASN	A	119	14.147	25.078	45.955	1.00	44.03	C
ATOM	834	CG	ASN	A	119	13.514	26.264	46.646	1.00	44.80	C
ATOM	835	OD1	ASN	A	119	14.207	27.129	47.197	1.00	46.56	O
ATOM	836	ND2	ASN	A	119	12.184	26.303	46.640	1.00	45.18	N
ATOM	837	N	ARG	A	120	16.380	23.533	44.337	1.00	42.37	N
ATOM	838	CA	ARG	A	120	16.778	22.278	43.716	1.00	41.69	C
ATOM	839	C	ARG	A	120	16.368	21.162	44.656	1.00	41.20	C
ATOM	840	O	ARG	A	120	16.577	21.267	45.859	1.00	41.46	O
ATOM	841	CB	ARG	A	120	18.299	22.239	43.504	1.00	41.70	C
ATOM	842	CG	ARG	A	120	18.827	21.002	42.754	1.00	40.96	C
ATOM	843	CD	ARG	A	120	20.351	20.897	42.683	1.00	39.92	C
ATOM	844	NE	ARG	A	120	21.001	21.327	43.920	1.00	39.89	N
ATOM	845	CZ	ARG	A	120	21.337	20.523	44.921	1.00	39.57	C
ATOM	846	NH1	ARG	A	120	21.084	19.217	44.856	1.00	39.43	N
ATOM	847	NH2	ARG	A	120	21.930	21.027	45.998	1.00	39.06	N
ATOM	848	N	GLU	A	121	15.769	20.109	44.115	1.00	40.44	N
ATOM	849	CA	GLU	A	121	15.380	18.956	44.908	1.00	39.92	C
ATOM	850	C	GLU	A	121	15.840	17.702	44.161	1.00	39.36	C
ATOM	851	O	GLU	A	121	15.716	17.617	42.942	1.00	39.20	O
ATOM	852	CB	GLU	A	121	13.866	18.950	45.170	1.00	39.97	C
ATOM	853	CG	GLU	A	121	13.369	17.698	45.876	1.00	40.42	C
ATOM	854	CD	GLU	A	121	12.136	17.928	46.731	1.00	41.22	C
ATOM	855	OE1	GLU	A	121	12.291	18.365	47.890	1.00	41.90	O
ATOM	856	OE2	GLU	A	121	11.016	17.645	46.259	1.00	42.21	O
ATOM	857	N	GLU	A	122	16.414	16.752	44.889	1.00	38.78	N
ATOM	858	CA	GLU	A	122	16.879	15.519	44.285	1.00	38.34	C
ATOM	859	C	GLU	A	122	15.760	14.523	44.374	1.00	38.13	C
ATOM	860	O	GLU	A	122	15.171	14.353	45.435	1.00	38.09	O
ATOM	861	CB	GLU	A	122	18.092	14.989	45.025	1.00	38.08	C
ATOM	862	CG	GLU	A	122	19.263	15.928	44.936	1.00	38.04	C
ATOM	863	CD	GLU	A	122	19.581	16.289	43.497	1.00	39.25	C
ATOM	864	OE1	GLU	A	122	19.763	15.364	42.667	1.00	38.89	O

ATOM	865	OE2	GLU	A	122	19.627	17.501	43.183	1.00	40.06	O
ATOM	866	N	MET	A	123	15.434	13.879	43.265	1.00	37.93	N
ATOM	867	CA	MET	A	123	14.384	12.885	43.312	1.00	37.84	C
ATOM	868	C	MET	A	123	14.536	11.819	42.242	1.00	37.41	C
ATOM	869	O	MET	A	123	15.418	11.876	41.387	1.00	37.04	O
ATOM	870	CB	MET	A	123	13.021	13.565	43.225	1.00	38.01	C
ATOM	871	CG	MET	A	123	12.798	14.318	41.946	1.00	39.05	C
ATOM	872	SD	MET	A	123	11.177	15.064	41.910	1.00	41.38	S
ATOM	873	CE	MET	A	123	11.489	16.589	42.818	1.00	41.39	C
ATOM	874	N	LYS	A	124	13.675	10.820	42.332	1.00	37.17	N
ATOM	875	CA	LYS	A	124	13.692	9.715	41.401	1.00	36.98	C
ATOM	876	C	LYS	A	124	12.802	10.098	40.238	1.00	36.81	C
ATOM	877	O	LYS	A	124	11.814	10.813	40.402	1.00	36.91	O
ATOM	878	CB	LYS	A	124	13.213	8.434	42.087	1.00	36.82	C
ATOM	879	CG	LYS	A	124	14.081	7.989	43.281	1.00	36.79	C
ATOM	880	CD	LYS	A	124	15.488	7.553	42.856	1.00	36.83	C
ATOM	881	CE	LYS	A	124	16.390	7.206	44.031	1.00	36.89	C
ATOM	882	NZ	LYS	A	124	17.822	7.412	43.680	1.00	36.99	N
ATOM	883	N	PHE	A	125	13.145	9.622	39.056	1.00	36.49	N
ATOM	884	CA	PHE	A	125	12.416	10.031	37.885	1.00	36.33	C
ATOM	885	C	PHE	A	125	10.926	9.861	38.074	1.00	36.45	C
ATOM	886	O	PHE	A	125	10.143	10.743	37.732	1.00	36.30	O
ATOM	887	CB	PHE	A	125	12.879	9.270	36.665	1.00	36.24	C
ATOM	888	CG	PHE	A	125	12.555	9.967	35.407	1.00	36.21	C
ATOM	889	CD1	PHE	A	125	13.403	10.936	34.924	1.00	35.98	C
ATOM	890	CD2	PHE	A	125	11.374	9.714	34.743	1.00	36.15	C
ATOM	891	CE1	PHE	A	125	13.108	11.616	33.794	1.00	36.07	C
ATOM	892	CE2	PHE	A	125	11.073	10.390	33.597	1.00	36.71	C
ATOM	893	CZ	PHE	A	125	11.943	11.350	33.120	1.00	36.73	C
ATOM	894	N	HIS	A	126	10.527	8.733	38.640	1.00	36.80	N
ATOM	895	CA	HIS	A	126	9.111	8.492	38.845	1.00	36.98	C
ATOM	896	C	HIS	A	126	8.547	9.560	39.772	1.00	37.20	C
ATOM	897	O	HIS	A	126	7.376	9.923	39.669	1.00	37.15	O
ATOM	898	CB	HIS	A	126	8.845	7.080	39.390	1.00	36.97	C
ATOM	899	CG	HIS	A	126	8.990	6.946	40.877	1.00	36.62	C
ATOM	900	ND1	HIS	A	126	7.935	7.115	41.747	1.00	35.91	N
ATOM	901	CD2	HIS	A	126	10.058	6.618	41.644	1.00	36.38	C
ATOM	902	CE1	HIS	A	126	8.352	6.921	42.986	1.00	35.63	C
ATOM	903	NE2	HIS	A	126	9.636	6.619	42.951	1.00	35.43	N
ATOM	904	N	GLU	A	127	9.383	10.082	40.663	1.00	37.34	N
ATOM	905	CA	GLU	A	127	8.912	11.087	41.599	1.00	37.55	C
ATOM	906	C	GLU	A	127	8.657	12.379	40.835	1.00	37.54	C
ATOM	907	O	GLU	A	127	7.657	13.061	41.049	1.00	37.36	O
ATOM	908	CB	GLU	A	127	9.914	11.308	42.736	1.00	37.65	C
ATOM	909	CG	GLU	A	127	10.155	10.091	43.617	1.00	37.89	C
ATOM	910	CD	GLU	A	127	11.113	10.383	44.764	1.00	38.14	C
ATOM	911	OE1	GLU	A	127	12.330	10.531	44.520	1.00	37.77	O
ATOM	912	OE2	GLU	A	127	10.649	10.471	45.917	1.00	38.86	O
ATOM	913	N	PHE	A	128	9.557	12.702	39.920	1.00	37.65	N
ATOM	914	CA	PHE	A	128	9.393	13.904	39.127	1.00	37.81	C
ATOM	915	C	PHE	A	128	8.090	13.798	38.350	1.00	37.95	C
ATOM	916	O	PHE	A	128	7.280	14.726	38.342	1.00	37.90	O
ATOM	917	CB	PHE	A	128	10.578	14.061	38.173	1.00	37.88	C
ATOM	918	CG	PHE	A	128	10.332	15.012	37.028	1.00	37.52	C
ATOM	919	CD1	PHE	A	128	10.088	16.355	37.258	1.00	36.94	C
ATOM	920	CD2	PHE	A	128	10.380	14.559	35.720	1.00	37.02	C
ATOM	921	CE1	PHE	A	128	9.876	17.219	36.203	1.00	37.09	C
ATOM	922	CE2	PHE	A	128	10.170	15.423	34.663	1.00	37.21	C
ATOM	923	CZ	PHE	A	128	9.917	16.748	34.901	1.00	37.15	C
ATOM	924	N	VAL	A	129	7.878	12.646	37.726	1.00	38.00	N
ATOM	925	CA	VAL	A	129	6.716	12.454	36.876	1.00	38.16	C

ATOM	926	C	VAL	A	129	5.442	12.637	37.671	1.00	38.24	C
ATOM	927	O	VAL	A	129	4.494	13.286	37.224	1.00	38.04	O
ATOM	928	CB	VAL	A	129	6.705	11.050	36.268	1.00	38.36	C
ATOM	929	CG1	VAL	A	129	5.398	10.810	35.508	1.00	38.41	C
ATOM	930	CG2	VAL	A	129	7.935	10.841	35.372	1.00	38.22	C
ATOM	931	N	GLU	A	130	5.430	12.038	38.854	1.00	38.42	N
ATOM	932	CA	GLU	A	130	4.289	12.115	39.745	1.00	38.54	C
ATOM	933	C	GLU	A	130	4.025	13.578	40.083	1.00	38.43	C
ATOM	934	O	GLU	A	130	2.898	14.056	39.960	1.00	38.29	O
ATOM	935	CB	GLU	A	130	4.547	11.265	40.997	1.00	38.69	C
ATOM	936	CG	GLU	A	130	4.404	9.763	40.745	1.00	39.15	C
ATOM	937	CD	GLU	A	130	5.160	8.897	41.740	1.00	39.75	C
ATOM	938	OE1	GLU	A	130	5.551	9.403	42.815	1.00	40.08	O
ATOM	939	OE2	GLU	A	130	5.355	7.697	41.441	1.00	40.15	O
ATOM	940	N	LYS	A	131	5.070	14.299	40.471	1.00	38.39	N
ATOM	941	CA	LYS	A	131	4.908	15.707	40.787	1.00	38.52	C
ATOM	942	C	LYS	A	131	4.290	16.431	39.598	1.00	38.61	C
ATOM	943	O	LYS	A	131	3.393	17.252	39.769	1.00	38.51	O
ATOM	944	CB	LYS	A	131	6.240	16.356	41.149	1.00	38.58	C
ATOM	945	CG	LYS	A	131	6.440	16.666	42.625	1.00	38.76	C
ATOM	946	CD	LYS	A	131	7.045	18.060	42.753	1.00	39.73	C
ATOM	947	CE	LYS	A	131	7.756	18.316	44.070	1.00	40.49	C
ATOM	948	NZ	LYS	A	131	8.117	19.772	44.197	1.00	40.68	N
ATOM	949	N	LEU	A	132	4.764	16.136	38.391	1.00	38.84	N
ATOM	950	CA	LEU	A	132	4.196	16.771	37.208	1.00	39.00	C
ATOM	951	C	LEU	A	132	2.726	16.429	37.113	1.00	39.11	C
ATOM	952	O	LEU	A	132	1.903	17.287	36.810	1.00	39.20	O
ATOM	953	CB	LEU	A	132	4.894	16.328	35.932	1.00	38.96	C
ATOM	954	CG	LEU	A	132	6.287	16.899	35.723	1.00	39.44	C
ATOM	955	CD1	LEU	A	132	6.870	16.389	34.429	1.00	39.81	C
ATOM	956	CD2	LEU	A	132	6.261	18.421	35.709	1.00	40.07	C
ATOM	957	N	GLN	A	133	2.396	15.170	37.367	1.00	39.22	N
ATOM	958	CA	GLN	A	133	1.005	14.758	37.335	1.00	39.38	C
ATOM	959	C	GLN	A	133	0.172	15.523	38.370	1.00	39.57	C
ATOM	960	O	GLN	A	133	-0.893	16.034	38.029	1.00	39.48	O
ATOM	961	CB	GLN	A	133	0.874	13.250	37.536	1.00	39.22	C
ATOM	962	N	ASP	A	134	0.656	15.631	39.611	1.00	39.91	N
ATOM	963	CA	ASP	A	134	-0.129	16.279	40.677	1.00	40.29	C
ATOM	964	C	ASP	A	134	-0.437	17.732	40.379	1.00	40.17	C
ATOM	965	O	ASP	A	134	-1.543	18.211	40.623	1.00	40.02	O
ATOM	966	CB	ASP	A	134	0.564	16.198	42.041	1.00	40.57	C
ATOM	967	CG	ASP	A	134	-0.348	16.660	43.182	1.00	41.99	C
ATOM	968	OD1	ASP	A	134	-1.301	17.421	42.910	1.00	43.76	O
ATOM	969	OD2	ASP	A	134	-0.212	16.312	44.378	1.00	44.05	O
ATOM	970	N	ILE	A	135	0.545	18.438	39.846	1.00	40.17	N
ATOM	971	CA	ILE	A	135	0.350	19.836	39.556	1.00	40.10	C
ATOM	972	C	ILE	A	135	-0.719	19.973	38.500	1.00	40.10	C
ATOM	973	O	ILE	A	135	-1.648	20.759	38.652	1.00	40.26	O
ATOM	974	CB	ILE	A	135	1.639	20.455	39.085	1.00	40.10	C
ATOM	975	CG1	ILE	A	135	2.612	20.560	40.261	1.00	40.11	C
ATOM	976	CG2	ILE	A	135	1.358	21.817	38.507	1.00	40.22	C
ATOM	977	CD1	ILE	A	135	4.056	20.688	39.842	1.00	40.17	C
ATOM	978	N	GLN	A	136	-0.596	19.198	37.431	1.00	40.02	N
ATOM	979	CA	GLN	A	136	-1.586	19.248	36.376	1.00	39.96	C
ATOM	980	C	GLN	A	136	-2.956	19.094	37.000	1.00	39.99	C
ATOM	981	O	GLN	A	136	-3.795	19.991	36.921	1.00	39.86	O
ATOM	982	CB	GLN	A	136	-1.350	18.144	35.361	1.00	40.00	C
ATOM	983	N	GLN	A	137	-3.158	17.972	37.679	1.00	40.18	N
ATOM	984	CA	GLN	A	137	-4.486	17.630	38.164	1.00	40.31	C
ATOM	985	C	GLN	A	137	-4.995	18.683	39.114	1.00	40.45	C
ATOM	986	O	GLN	A	137	-6.170	19.046	39.068	1.00	40.75	O

ATOM	987	CB	GLN	A	137	-4.502	16.247	38.829	1.00	40.24	C
ATOM	988	N	ARG	A	138	-4.116	19.200	39.959	1.00	40.45	N
ATOM	989	CA	ARG	A	138	-4.560	20.159	40.956	1.00	40.45	C
ATOM	990	C	ARG	A	138	-4.568	21.588	40.419	1.00	40.54	C
ATOM	991	O	ARG	A	138	-4.678	22.541	41.188	1.00	40.67	O
ATOM	992	CB	ARG	A	138	-3.692	20.069	42.207	1.00	40.43	C
ATOM	993	CG	ARG	A	138	-2.399	20.868	42.161	1.00	40.03	C
ATOM	994	CD	ARG	A	138	-1.541	20.546	43.346	1.00	39.94	C
ATOM	995	NE	ARG	A	138	-0.431	21.458	43.580	1.00	39.71	N
ATOM	996	CZ	ARG	A	138	0.842	21.098	43.498	1.00	40.13	C
ATOM	997	NH1	ARG	A	138	1.164	19.857	43.150	1.00	40.53	N
ATOM	998	NH2	ARG	A	138	1.800	21.978	43.747	1.00	40.33	N
ATOM	999	N	GLY	A	139	-4.471	21.742	39.102	1.00	40.47	N
ATOM	1000	CA	GLY	A	139	-4.418	23.066	38.508	1.00	40.46	C
ATOM	1001	C	GLY	A	139	-3.432	24.021	39.176	1.00	40.45	C
ATOM	1002	O	GLY	A	139	-3.616	25.235	39.110	1.00	40.46	O
ATOM	1003	N	GLY	A	140	-2.370	23.490	39.780	1.00	40.47	N
ATOM	1004	CA	GLY	A	140	-1.425	24.314	40.519	1.00	40.45	C
ATOM	1005	C	GLY	A	140	-0.695	25.272	39.602	1.00	40.40	C
ATOM	1006	O	GLY	A	140	-0.784	25.116	38.383	1.00	40.79	O
ATOM	1007	N	GLU	A	141	-0.015	26.273	40.167	1.00	40.10	N
ATOM	1008	CA	GLU	A	141	0.807	27.195	39.371	1.00	40.01	C
ATOM	1009	C	GLU	A	141	2.305	26.854	39.458	1.00	39.64	C
ATOM	1010	O	GLU	A	141	3.112	27.386	38.692	1.00	39.52	O
ATOM	1011	CB	GLU	A	141	0.589	28.660	39.796	1.00	40.20	C
ATOM	1012	CG	GLU	A	141	-0.711	29.298	39.315	1.00	41.00	C
ATOM	1013	CD	GLU	A	141	-0.845	29.307	37.800	1.00	42.10	C
ATOM	1014	OE1	GLU	A	141	0.194	29.181	37.113	1.00	41.99	O
ATOM	1015	OE2	GLU	A	141	-1.990	29.437	37.300	1.00	43.15	O
ATOM	1016	N	GLU	A	142	2.676	25.977	40.392	1.00	39.25	N
ATOM	1017	CA	GLU	A	142	4.075	25.589	40.563	1.00	38.95	C
ATOM	1018	C	GLU	A	142	4.613	25.162	39.210	1.00	38.74	C
ATOM	1019	O	GLU	A	142	3.859	24.694	38.351	1.00	38.77	O
ATOM	1020	CB	GLU	A	142	4.220	24.433	41.577	1.00	38.95	C
ATOM	1021	CG	GLU	A	142	5.658	24.138	42.019	1.00	38.90	C
ATOM	1022	CD	GLU	A	142	5.797	22.950	42.971	1.00	38.93	C
ATOM	1023	OE1	GLU	A	142	4.836	22.164	43.130	1.00	39.58	O
ATOM	1024	OE2	GLU	A	142	6.888	22.799	43.565	1.00	38.74	O
ATOM	1025	N	ARG	A	143	5.914	25.332	39.020	1.00	38.39	N
ATOM	1026	CA	ARG	A	143	6.569	24.908	37.796	1.00	38.25	C
ATOM	1027	C	ARG	A	143	7.809	24.122	38.122	1.00	38.14	C
ATOM	1028	O	ARG	A	143	8.512	24.450	39.078	1.00	38.46	O
ATOM	1029	CB	ARG	A	143	7.046	26.107	37.003	1.00	38.26	C
ATOM	1030	CG	ARG	A	143	5.986	26.945	36.398	1.00	37.89	C
ATOM	1031	CD	ARG	A	143	6.602	28.026	35.555	1.00	37.35	C
ATOM	1032	NE	ARG	A	143	5.610	28.814	34.849	1.00	36.66	N
ATOM	1033	CZ	ARG	A	143	5.316	28.668	33.573	1.00	36.40	C
ATOM	1034	NH1	ARG	A	143	5.933	27.755	32.831	1.00	36.41	N
ATOM	1035	NH2	ARG	A	143	4.397	29.443	33.034	1.00	37.02	N
ATOM	1036	N	LEU	A	144	8.116	23.124	37.304	1.00	37.71	N
ATOM	1037	CA	LEU	A	144	9.319	22.349	37.517	1.00	37.57	C
ATOM	1038	C	LEU	A	144	10.181	22.319	36.294	1.00	37.33	C
ATOM	1039	O	LEU	A	144	9.710	22.466	35.165	1.00	37.31	O
ATOM	1040	CB	LEU	A	144	8.979	20.909	37.858	1.00	37.69	C
ATOM	1041	CG	LEU	A	144	7.981	20.736	38.993	1.00	38.31	C
ATOM	1042	CD1	LEU	A	144	7.612	19.255	39.165	1.00	39.02	C
ATOM	1043	CD2	LEU	A	144	8.551	21.327	40.263	1.00	37.91	C
ATOM	1044	N	TYR	A	145	11.460	22.098	36.528	1.00	37.01	N
ATOM	1045	CA	TYR	A	145	12.379	21.922	35.437	1.00	36.87	C
ATOM	1046	C	TYR	A	145	13.397	20.886	35.871	1.00	36.85	C
ATOM	1047	O	TYR	A	145	14.146	21.093	36.819	1.00	36.51	O

ATOM	1048	CB	TYR	A	145	13.043	23.255	35.110	1.00	36.92	
ATOM	1049	CG	TYR	A	145	13.505	23.446	33.686	1.00	36.38	C
ATOM	1050	CD1	TYR	A	145	13.845	22.383	32.875	1.00	36.31	C
ATOM	1051	CD2	TYR	A	145	13.619	24.716	33.163	1.00	36.28	C
ATOM	1052	CE1	TYR	A	145	14.278	22.590	31.582	1.00	36.21	C
ATOM	1053	CE2	TYR	A	145	14.047	24.923	31.880	1.00	35.94	C
ATOM	1054	CZ	TYR	A	145	14.374	23.862	31.093	1.00	35.54	C
ATOM	1055	OH	TYR	A	145	14.789	24.081	29.799	1.00	36.13	O
ATOM	1056	N	LEU	A	146	13.378	19.736	35.218	1.00	37.06	N
ATOM	1057	CA	LEU	A	146	14.396	18.742	35.477	1.00	37.49	C
ATOM	1058	C	LEU	A	146	15.628	19.029	34.629	1.00	37.55	C
ATOM	1059	O	LEU	A	146	15.532	19.241	33.427	1.00	37.31	O
ATOM	1060	CB	LEU	A	146	13.877	17.341	35.173	1.00	37.69	C
ATOM	1061	CG	LEU	A	146	14.919	16.233	35.349	1.00	37.95	C
ATOM	1062	CD1	LEU	A	146	14.234	14.916	35.567	1.00	37.66	C
ATOM	1063	CD2	LEU	A	146	15.849	16.133	34.151	1.00	39.00	C
ATOM	1064	N	GLN	A	147	16.786	18.993	35.269	1.00	37.84	N
ATOM	1065	CA	GLN	A	147	18.053	19.259	34.617	1.00	38.09	C
ATOM	1066	C	GLN	A	147	19.088	18.397	35.306	1.00	37.92	C
ATOM	1067	O	GLN	A	147	19.453	18.661	36.443	1.00	37.87	O
ATOM	1068	CB	GLN	A	147	18.421	20.725	34.773	1.00	38.37	C
ATOM	1069	CG	GLN	A	147	17.331	21.686	34.362	1.00	39.01	C
ATOM	1070	CD	GLN	A	147	17.867	23.064	34.191	1.00	39.92	C
ATOM	1071	OE1	GLN	A	147	17.149	23.955	33.772	1.00	42.51	O
ATOM	1072	NE2	GLN	A	147	19.137	23.253	34.520	1.00	40.24	N
ATOM	1073	N	GLN	A	148	19.576	17.385	34.601	1.00	37.81	N
ATOM	1074	CA	GLN	A	148	20.433	16.377	35.196	1.00	37.57	C
ATOM	1075	C	GLN	A	148	21.325	15.744	34.176	1.00	37.68	C
ATOM	1076	O	GLN	A	148	20.876	15.266	33.146	1.00	37.30	O
ATOM	1077	CB	GLN	A	148	19.576	15.261	35.773	1.00	37.61	C
ATOM	1078	CG	GLN	A	148	20.336	13.971	36.068	1.00	36.79	C
ATOM	1079	CD	GLN	A	148	21.398	14.159	37.117	1.00	35.97	C
ATOM	1080	OE1	GLN	A	148	21.162	14.814	38.135	1.00	35.11	O
ATOM	1081	NE2	GLN	A	148	22.577	13.593	36.875	1.00	35.03	N
ATOM	1082	N	THR	A	149	22.596	15.698	34.503	1.00	38.15	N
ATOM	1083	CA	THR	A	149	23.574	15.130	33.622	1.00	38.57	C
ATOM	1084	C	THR	A	149	23.354	13.647	33.506	1.00	38.68	C
ATOM	1085	O	THR	A	149	23.154	12.973	34.509	1.00	39.07	O
ATOM	1086	CB	THR	A	149	24.936	15.410	34.214	1.00	38.75	C
ATOM	1087	OG1	THR	A	149	25.206	16.807	34.088	1.00	38.86	O
ATOM	1088	CG2	THR	A	149	26.036	14.759	33.418	1.00	39.39	C
ATOM	1089	N	LEU	A	150	23.393	13.144	32.279	1.00	38.87	N
ATOM	1090	CA	LEU	A	150	23.259	11.722	32.021	1.00	39.24	C
ATOM	1091	C	LEU	A	150	24.498	10.979	32.507	1.00	39.20	C
ATOM	1092	O	LEU	A	150	25.602	11.317	32.101	1.00	39.20	O
ATOM	1093	CB	LEU	A	150	23.146	11.465	30.519	1.00	39.46	C
ATOM	1094	CG	LEU	A	150	21.938	11.942	29.722	1.00	40.52	C
ATOM	1095	CD1	LEU	A	150	22.077	11.563	28.237	1.00	41.06	C
ATOM	1096	CD2	LEU	A	150	20.712	11.308	30.303	1.00	41.66	C
ATOM	1097	N	ASN	A	151	24.315	9.960	33.343	1.00	39.19	N
ATOM	1098	CA	ASN	A	151	25.436	9.171	33.861	1.00	39.46	C
ATOM	1099	C	ASN	A	151	25.193	7.642	33.883	1.00	39.43	C
ATOM	1100	O	ASN	A	151	24.198	7.162	33.333	1.00	39.15	O
ATOM	1101	CB	ASN	A	151	25.731	9.652	35.265	1.00	39.54	C
ATOM	1102	CG	ASN	A	151	24.511	9.588	36.150	1.00	40.08	C
ATOM	1103	OD1	ASN	A	151	23.874	8.533	36.283	1.00	39.68	O
ATOM	1104	ND2	ASN	A	151	24.158	10.725	36.751	1.00	40.83	N
ATOM	1105	N	ASP	A	152	26.075	6.892	34.556	1.00	39.43	N
ATOM	1106	CA	ASP	A	152	26.025	5.413	34.543	1.00	39.58	C
ATOM	1107	C	ASP	A	152	24.761	4.787	35.083	1.00	39.17	C
ATOM	1108	O	ASP	A	152	24.477	3.634	34.776	1.00	39.29	O

ATOM	1109	CB	ASP	A	152	27.138	4.756	35.385	1.00	39.92	C
ATOM	1110	CG	ASP	A	152	28.489	5.382	35.198	1.00	41.53	O
ATOM	1111	OD1	ASP	A	152	28.722	6.005	34.134	1.00	45.10	C
ATOM	1112	OD2	ASP	A	152	29.378	5.294	36.076	1.00	42.02	O
ATOM	1113	N	THR	A	153	24.008	5.493	35.909	1.00	38.78	N
ATOM	1114	CA	THR	A	153	22.881	4.830	36.542	1.00	38.50	C
ATOM	1115	C	THR	A	153	21.757	4.573	35.573	1.00	38.26	C
ATOM	1116	O	THR	A	153	20.842	3.838	35.882	1.00	38.29	O
ATOM	1117	CB	THR	A	153	22.329	5.624	37.733	1.00	38.46	C
ATOM	1118	OG1	THR	A	153	21.622	6.780	37.269	1.00	39.25	O
ATOM	1119	CG2	THR	A	153	23.445	6.161	38.611	1.00	38.28	C
ATOM	1120	N	VAL	A	154	21.804	5.165	34.396	1.00	38.28	N
ATOM	1121	CA	VAL	A	154	20.687	4.977	33.491	1.00	38.29	C
ATOM	1122	C	VAL	A	154	20.602	3.530	33.067	1.00	38.17	C
ATOM	1123	O	VAL	A	154	21.597	2.805	33.092	1.00	38.10	O
ATOM	1124	CB	VAL	A	154	20.786	5.838	32.251	1.00	38.15	C
ATOM	1125	CG1	VAL	A	154	20.815	7.291	32.649	1.00	38.22	C
ATOM	1126	CG2	VAL	A	154	22.007	5.437	31.443	1.00	38.40	C
ATOM	1127	N	GLY	A	155	19.395	3.132	32.681	1.00	38.05	N
ATOM	1128	CA	GLY	A	155	19.114	1.779	32.258	1.00	38.07	C
ATOM	1129	C	GLY	A	155	19.609	1.358	30.894	1.00	38.17	C
ATOM	1130	O	GLY	A	155	20.092	2.153	30.081	1.00	38.44	O
ATOM	1131	N	ARG	A	156	19.417	0.073	30.642	1.00	38.16	N
ATOM	1132	CA	ARG	A	156	19.932	-0.588	29.461	1.00	38.16	C
ATOM	1133	C	ARG	A	156	19.414	0.065	28.210	1.00	38.01	C
ATOM	1134	O	ARG	A	156	20.185	0.490	27.346	1.00	38.04	O
ATOM	1135	CB	ARG	A	156	19.554	-2.080	29.489	1.00	38.18	C
ATOM	1136	N	LYS	A	157	18.102	0.178	28.118	1.00	37.83	N
ATOM	1137	CA	LYS	A	157	17.545	0.713	26.905	1.00	37.67	C
ATOM	1138	C	LYS	A	157	18.144	2.092	26.679	1.00	37.77	C
ATOM	1139	O	LYS	A	157	18.579	2.386	25.571	1.00	37.84	O
ATOM	1140	CB	LYS	A	157	16.014	0.725	26.936	1.00	37.62	C
ATOM	1141	CG	LYS	A	157	15.379	-0.624	26.550	1.00	36.92	C
ATOM	1142	N	ILE	A	158	18.247	2.911	27.728	1.00	37.82	N
ATOM	1143	CA	ILE	A	158	18.699	4.287	27.526	1.00	37.81	C
ATOM	1144	C	ILE	A	158	20.096	4.252	27.008	1.00	37.78	C
ATOM	1145	O	ILE	A	158	20.440	4.942	26.058	1.00	37.52	O
ATOM	1146	CB	ILE	A	158	18.666	5.109	28.802	1.00	37.90	C
ATOM	1147	CG1	ILE	A	158	17.233	5.283	29.280	1.00	37.87	C
ATOM	1148	CG2	ILE	A	158	19.281	6.477	28.538	1.00	37.57	C
ATOM	1149	CD1	ILE	A	158	16.383	6.034	28.318	1.00	37.95	C
ATOM	1150	N	VAL	A	159	20.899	3.418	27.637	1.00	37.89	N
ATOM	1151	CA	VAL	A	159	22.263	3.265	27.211	1.00	38.06	C
ATOM	1152	C	VAL	A	159	22.255	2.906	25.725	1.00	38.07	C
ATOM	1153	O	VAL	A	159	23.025	3.462	24.955	1.00	37.93	O
ATOM	1154	CB	VAL	A	159	22.987	2.196	28.039	1.00	38.12	C
ATOM	1155	CG1	VAL	A	159	24.208	1.740	27.315	1.00	38.65	C
ATOM	1156	CG2	VAL	A	159	23.368	2.734	29.410	1.00	37.65	C
ATOM	1157	N	MET	A	160	21.385	1.984	25.318	1.00	38.30	N
ATOM	1158	CA	MET	A	160	21.264	1.658	23.894	1.00	38.71	C
ATOM	1159	C	MET	A	160	21.024	2.923	23.084	1.00	38.35	C
ATOM	1160	O	MET	A	160	21.742	3.219	22.137	1.00	38.42	O
ATOM	1161	CB	MET	A	160	20.115	0.682	23.636	1.00	38.93	C
ATOM	1162	CG	MET	A	160	20.431	-0.690	24.087	1.00	40.64	C
ATOM	1163	SD	MET	A	160	21.840	-1.293	23.180	1.00	44.02	S
ATOM	1164	CE	MET	A	160	20.982	-1.709	21.617	1.00	44.00	C
ATOM	1165	N	ASP	A	161	19.997	3.665	23.458	1.00	37.94	N
ATOM	1166	CA	ASP	A	161	19.702	4.902	22.773	1.00	37.82	C
ATOM	1167	C	ASP	A	161	20.955	5.788	22.684	1.00	37.50	C
ATOM	1168	O	ASP	A	161	21.395	6.154	21.596	1.00	37.66	O
ATOM	1169	CB	ASP	A	161	18.591	5.650	23.511	1.00	37.84	C

ATOM	1231	C	ASN	A	168	30.447	11.229	13.590	1.00	32.97	C
ATOM	1232	O	ASN	A	168	31.541	11.747	13.390	1.00	33.15	O
ATOM	1233	CB	ASN	A	168	29.729	9.064	12.749	1.00	32.68	C
ATOM	1234	CG	ASN	A	168	30.594	9.147	11.507	1.00	32.86	C
ATOM	1235	OD1	ASN	A	168	31.797	8.914	11.574	1.00	35.02	O
ATOM	1236	ND2	ASN	A	168	29.989	9.473	10.368	1.00	31.39	N
ATOM	1237	N	TRP	A	169	29.284	11.884	13.518	1.00	33.38	N
ATOM	1238	CA	TRP	A	169	29.200	13.279	13.107	1.00	33.28	C
ATOM	1239	C	TRP	A	169	29.711	14.237	14.149	1.00	33.33	C
ATOM	1240	O	TRP	A	169	30.406	15.188	13.834	1.00	33.43	O
ATOM	1241	CB	TRP	A	169	27.756	13.673	12.765	1.00	33.43	C
ATOM	1242	CG	TRP	A	169	27.618	15.134	12.412	1.00	33.11	C
ATOM	1243	CD1	TRP	A	169	27.814	15.698	11.191	1.00	33.49	C
ATOM	1244	CD2	TRP	A	169	27.293	16.205	13.294	1.00	32.94	C
ATOM	1245	NE1	TRP	A	169	27.629	17.056	11.251	1.00	33.21	N
ATOM	1246	CE2	TRP	A	169	27.306	17.397	12.533	1.00	33.42	C
ATOM	1247	CE3	TRP	A	169	26.994	16.284	14.655	1.00	33.15	C
ATOM	1248	CZ2	TRP	A	169	27.026	18.648	13.081	1.00	33.66	C
ATOM	1249	CZ3	TRP	A	169	26.719	17.526	15.204	1.00	34.02	C
ATOM	1250	CH2	TRP	A	169	26.733	18.695	14.413	1.00	34.21	C
ATOM	1251	N	ILE	A	170	29.369	14.021	15.398	1.00	33.54	N
ATOM	1252	CA	ILE	A	170	29.739	15.024	16.368	1.00	33.92	C
ATOM	1253	C	ILE	A	170	31.225	14.935	16.713	1.00	34.07	C
ATOM	1254	O	ILE	A	170	31.855	15.939	16.988	1.00	34.23	O
ATOM	1255	CB	ILE	A	170	28.832	14.947	17.591	1.00	33.84	C
ATOM	1256	CG1	ILE	A	170	28.803	16.288	18.306	1.00	34.22	C
ATOM	1257	CG2	ILE	A	170	29.295	13.858	18.499	1.00	34.45	C
ATOM	1258	CD1	ILE	A	170	27.869	16.309	19.478	1.00	34.73	C
ATOM	1259	N	ASN	A	171	31.801	13.743	16.650	1.00	34.41	N
ATOM	1260	CA	ASN	A	171	33.222	13.593	16.950	1.00	34.46	C
ATOM	1261	C	ASN	A	171	34.055	14.322	15.918	1.00	34.56	C
ATOM	1262	O	ASN	A	171	35.055	14.949	16.260	1.00	34.66	O
ATOM	1263	CB	ASN	A	171	33.632	12.116	16.996	1.00	34.47	C
ATOM	1264	CG	ASN	A	171	32.969	11.356	18.128	1.00	33.99	C
ATOM	1265	OD1	ASN	A	171	32.391	11.951	19.031	1.00	35.35	O
ATOM	1266	ND2	ASN	A	171	33.040	10.036	18.077	1.00	32.53	N
ATOM	1267	N	LYS	A	172	33.652	14.217	14.654	1.00	34.75	N
ATOM	1268	CA	LYS	A	172	34.337	14.917	13.584	1.00	35.18	C
ATOM	1269	C	LYS	A	172	34.261	16.415	13.907	1.00	35.29	C
ATOM	1270	O	LYS	A	172	35.248	17.133	13.800	1.00	34.89	O
ATOM	1271	CB	LYS	A	172	33.711	14.600	12.216	1.00	35.26	C
ATOM	1272	CG	LYS	A	172	34.200	13.299	11.587	1.00	36.55	C
ATOM	1273	CD	LYS	A	172	33.873	13.158	10.090	1.00	38.20	C
ATOM	1274	CE	LYS	A	172	34.512	11.872	9.537	1.00	39.82	C
ATOM	1275	NZ	LYS	A	172	34.369	11.661	8.050	1.00	41.20	N
ATOM	1276	N	GLN	A	173	33.093	16.875	14.339	1.00	35.70	N
ATOM	1277	CA	GLN	A	173	32.931	18.272	14.699	1.00	36.20	C
ATOM	1278	C	GLN	A	173	33.929	18.633	15.771	1.00	36.47	C
ATOM	1279	O	GLN	A	173	34.605	19.656	15.686	1.00	36.97	O
ATOM	1280	CB	GLN	A	173	31.534	18.555	15.231	1.00	36.19	C
ATOM	1281	CG	GLN	A	173	30.473	18.637	14.167	1.00	36.97	C
ATOM	1282	CD	GLN	A	173	30.765	19.720	13.157	1.00	37.19	C
ATOM	1283	OE1	GLN	A	173	31.198	20.808	13.531	1.00	38.68	O
ATOM	1284	NE2	GLN	A	173	30.551	19.425	11.879	1.00	35.94	N
ATOM	1285	N	GLN	A	174	34.000	17.806	16.802	1.00	36.38	N
ATOM	1286	CA	GLN	A	174	34.945	18.041	17.866	1.00	36.42	C
ATOM	1287	C	GLN	A	174	36.360	18.102	17.283	1.00	36.56	C
ATOM	1288	O	GLN	A	174	37.161	18.945	17.672	1.00	36.42	O
ATOM	1289	CB	GLN	A	174	34.815	16.937	18.912	1.00	36.33	C
ATOM	1290	CG	GLN	A	174	35.896	16.908	19.963	1.00	36.13	C
ATOM	1291	CD	GLN	A	174	35.785	15.676	20.815	1.00	36.16	C

ATOM	1292	OE1	GLN	A	174	35.415	14.613	20.314	1.00	37.84	O
ATOM	1293	NE2	GLN	A	174	36.082	15.803	22.103	1.00	34.98	N
ATOM	1294	N	GLY	A	175	36.649	17.231	16.322	1.00	36.76	N
ATOM	1295	CA	GLY	A	175	37.976	17.158	15.750	1.00	36.97	C
ATOM	1296	C	GLY	A	175	38.302	18.356	14.892	1.00	37.21	C
ATOM	1297	O	GLY	A	175	39.336	18.993	15.070	1.00	37.38	O
ATOM	1298	N	LYS	A	176	37.406	18.682	13.971	1.00	37.44	N
ATOM	1299	CA	LYS	A	176	37.634	19.793	13.064	1.00	37.56	C
ATOM	1300	C	LYS	A	176	37.898	21.090	13.822	1.00	37.47	C
ATOM	1301	O	LYS	A	176	38.713	21.907	13.400	1.00	37.47	O
ATOM	1302	CB	LYS	A	176	36.423	20.014	12.161	1.00	37.71	C
ATOM	1303	CG	LYS	A	176	36.134	18.905	11.163	1.00	38.70	C
ATOM	1304	CD	LYS	A	176	34.960	19.284	10.279	1.00	39.76	C
ATOM	1305	CE	LYS	A	176	34.064	18.092	10.037	1.00	41.48	C
ATOM	1306	NZ	LYS	A	176	32.663	18.481	9.690	1.00	43.43	N
ATOM	1307	N	ARG	A	177	37.212	21.283	14.938	1.00	37.10	N
ATOM	1308	CA	ARG	A	177	37.325	22.544	15.643	1.00	37.08	C
ATOM	1309	C	ARG	A	177	38.352	22.495	16.763	1.00	36.85	C
ATOM	1310	O	ARG	A	177	38.445	23.420	17.564	1.00	36.21	O
ATOM	1311	CB	ARG	A	177	35.980	22.969	16.228	1.00	37.21	C
ATOM	1312	CG	ARG	A	177	34.807	22.773	15.326	1.00	37.50	C
ATOM	1313	CD	ARG	A	177	34.802	23.591	14.053	1.00	38.49	C
ATOM	1314	NE	ARG	A	177	33.750	23.066	13.185	1.00	39.56	N
ATOM	1315	CZ	ARG	A	177	33.884	22.803	11.891	1.00	41.17	C
ATOM	1316	NH1	ARG	A	177	35.026	23.050	11.254	1.00	41.34	N
ATOM	1317	NH2	ARG	A	177	32.855	22.304	11.219	1.00	41.89	N
ATOM	1318	N	GLY	A	178	39.088	21.401	16.859	1.00	36.70	N
ATOM	1319	CA	GLY	A	178	40.131	21.337	17.854	1.00	36.74	C
ATOM	1320	C	GLY	A	178	39.592	21.509	19.249	1.00	36.76	C
ATOM	1321	O	GLY	A	178	40.340	21.817	20.171	1.00	37.18	O
ATOM	1322	N	TRP	A	179	38.293	21.331	19.420	1.00	36.75	N
ATOM	1323	CA	TRP	A	179	37.722	21.439	20.745	1.00	36.66	C
ATOM	1324	C	TRP	A	179	38.309	20.400	21.679	1.00	36.60	C
ATOM	1325	O	TRP	A	179	38.980	19.451	21.260	1.00	35.99	O
ATOM	1326	CB	TRP	A	179	36.222	21.216	20.708	1.00	36.79	C
ATOM	1327	CG	TRP	A	179	35.454	22.258	20.038	1.00	36.51	C
ATOM	1328	CD1	TRP	A	179	35.889	23.470	19.620	1.00	36.67	C
ATOM	1329	CD2	TRP	A	179	34.081	22.180	19.692	1.00	37.40	C
ATOM	1330	NE1	TRP	A	179	34.860	24.168	19.035	1.00	36.91	N
ATOM	1331	CE2	TRP	A	179	33.735	23.390	19.062	1.00	37.43	C
ATOM	1332	CE3	TRP	A	179	33.099	21.198	19.840	1.00	36.64	C
ATOM	1333	CZ2	TRP	A	179	32.456	23.641	18.583	1.00	36.83	C
ATOM	1334	CZ3	TRP	A	179	31.840	21.450	19.365	1.00	36.84	C
ATOM	1335	CH2	TRP	A	179	31.524	22.661	18.744	1.00	36.02	C
ATOM	1336	N	GLY	A	180	38.022	20.595	22.958	1.00	36.66	N
ATOM	1337	CA	GLY	A	180	38.444	19.681	23.992	1.00	36.96	C
ATOM	1338	C	GLY	A	180	37.456	18.548	24.131	1.00	37.25	C
ATOM	1339	O	GLY	A	180	36.698	18.285	23.204	1.00	37.69	O
ATOM	1340	N	GLN	A	181	37.445	17.891	25.288	1.00	37.46	N
ATOM	1341	CA	GLN	A	181	36.581	16.734	25.507	1.00	37.81	C
ATOM	1342	C	GLN	A	181	35.136	17.088	25.802	1.00	37.40	C
ATOM	1343	O	GLN	A	181	34.825	18.177	26.273	1.00	37.59	O
ATOM	1344	CB	GLN	A	181	37.082	15.881	26.680	1.00	38.18	C
ATOM	1345	CG	GLN	A	181	36.684	16.433	28.065	1.00	40.32	C
ATOM	1346	CD	GLN	A	181	36.956	15.470	29.231	1.00	43.44	C
ATOM	1347	OE1	GLN	A	181	38.004	14.802	29.284	1.00	44.84	O
ATOM	1348	NE2	GLN	A	181	36.007	15.406	30.173	1.00	44.80	N
ATOM	1349	N	LEU	A	182	34.266	16.130	25.510	1.00	37.00	N
ATOM	1350	CA	LEU	A	182	32.866	16.181	25.878	1.00	36.47	C
ATOM	1351	C	LEU	A	182	32.856	16.224	27.385	1.00	35.97	C
ATOM	1352	O	LEU	A	182	33.345	15.298	28.009	1.00	35.98	O

ATOM	1353	CB	LEU A 182	32.208	14.865	25.468	1.00	36.38	C
ATOM	1354	CG	LEU A 182	30.691	14.707	25.336	1.00	36.76	C
ATOM	1355	CD1	LEU A 182	30.339	13.254	25.433	1.00	36.70	C
ATOM	1356	CD2	LEU A 182	29.917	15.417	26.376	1.00	37.66	C
ATOM	1357	N	THR A 183	32.337	17.270	28.005	1.00	35.50	N
ATOM	1358	CA	THR A 183	32.310	17.230	29.457	1.00	35.07	C
ATOM	1359	C	THR A 183	31.008	16.669	29.929	1.00	34.39	C
ATOM	1360	O	THR A 183	30.944	16.097	31.000	1.00	34.13	O
ATOM	1361	CB	THR A 183	32.528	18.599	30.105	1.00	35.07	C
ATOM	1362	OG1	THR A 183	31.402	19.444	29.883	1.00	35.21	O
ATOM	1363	CG2	THR A 183	33.652	19.310	29.453	1.00	35.76	C
ATOM	1364	N	SER A 184	29.947	16.848	29.163	1.00	33.83	N
ATOM	1365	CA	SER A 184	28.710	16.288	29.626	1.00	33.19	C
ATOM	1366	C	SER A 184	27.563	16.459	28.684	1.00	32.72	C
ATOM	1367	O	SER A 184	27.643	17.172	27.699	1.00	32.59	O
ATOM	1368	CB	SER A 184	28.363	16.954	30.949	1.00	33.18	C
ATOM	1369	OG	SER A 184	28.349	18.347	30.788	1.00	31.82	O
ATOM	1370	N	ASN A 185	26.484	15.776	29.013	1.00	32.44	N
ATOM	1371	CA	ASN A 185	25.259	15.893	28.277	1.00	32.59	C
ATOM	1372	C	ASN A 185	24.218	16.117	29.319	1.00	33.00	C
ATOM	1373	O	ASN A 185	24.027	15.285	30.184	1.00	32.72	O
ATOM	1374	CB	ASN A 185	24.917	14.619	27.520	1.00	32.46	C
ATOM	1375	CG	ASN A 185	25.850	14.351	26.369	1.00	31.69	C
ATOM	1376	OD1	ASN A 185	26.661	13.436	26.434	1.00	32.60	O
ATOM	1377	ND2	ASN A 185	25.725	15.121	25.297	1.00	29.86	N
ATOM	1378	N	LEU A 186	23.549	17.252	29.258	1.00	33.80	N
ATOM	1379	CA	LEU A 186	22.537	17.536	30.241	1.00	34.36	C
ATOM	1380	C	LEU A 186	21.199	17.117	29.684	1.00	34.78	C
ATOM	1381	O	LEU A 186	20.903	17.380	28.512	1.00	35.02	O
ATOM	1382	CB	LEU A 186	22.500	19.029	30.539	1.00	34.34	C
ATOM	1383	CG	LEU A 186	21.564	19.433	31.677	1.00	35.25	C
ATOM	1384	CD1	LEU A 186	22.122	18.827	32.930	1.00	35.58	C
ATOM	1385	CD2	LEU A 186	21.402	20.956	31.845	1.00	35.43	C
ATOM	1386	N	LEU A 187	20.393	16.456	30.507	1.00	34.95	N
ATOM	1387	CA	LEU A 187	19.022	16.189	30.120	1.00	35.27	C
ATOM	1388	C	LEU A 187	18.117	17.260	30.735	1.00	35.48	C
ATOM	1389	O	LEU A 187	18.105	17.449	31.951	1.00	35.40	O
ATOM	1390	CB	LEU A 187	18.593	14.805	30.563	1.00	35.33	C
ATOM	1391	CG	LEU A 187	17.105	14.481	30.399	1.00	36.09	C
ATOM	1392	CD1	LEU A 187	16.624	14.525	28.969	1.00	36.40	C
ATOM	1393	CD2	LEU A 187	16.850	13.114	30.940	1.00	36.96	C
ATOM	1394	N	LEU A 188	17.380	17.977	29.889	1.00	35.70	N
ATOM	1395	CA	LEU A 188	16.466	19.005	30.362	1.00	35.89	C
ATOM	1396	C	LEU A 188	15.047	18.668	29.972	1.00	36.32	C
ATOM	1397	O	LEU A 188	14.738	18.548	28.792	1.00	36.89	O
ATOM	1398	CB	LEU A 188	16.795	20.362	29.771	1.00	35.79	C
ATOM	1399	CG	LEU A 188	18.210	20.889	29.937	1.00	35.66	C
ATOM	1400	CD1	LEU A 188	18.976	20.745	28.668	1.00	36.04	C
ATOM	1401	CD2	LEU A 188	18.125	22.328	30.284	1.00	35.23	C
ATOM	1402	N	ILE A 189	14.184	18.514	30.967	1.00	36.45	N
ATOM	1403	CA	ILE A 189	12.782	18.254	30.720	1.00	36.25	C
ATOM	1404	C	ILE A 189	12.105	19.353	31.472	1.00	35.88	C
ATOM	1405	O	ILE A 189	12.388	19.542	32.651	1.00	35.75	O
ATOM	1406	CB	ILE A 189	12.368	16.916	31.274	1.00	36.47	C
ATOM	1407	CG1	ILE A 189	13.126	15.813	30.549	1.00	36.63	C
ATOM	1408	CG2	ILE A 189	10.865	16.748	31.143	1.00	36.55	C
ATOM	1409	CD1	ILE A 189	12.884	14.440	31.129	1.00	36.11	C
ATOM	1410	N	GLY A 190	11.221	20.071	30.789	1.00	35.53	N
ATOM	1411	CA	GLY A 190	10.614	21.257	31.344	1.00	35.37	C
ATOM	1412	C	GLY A 190	9.184	21.420	30.921	1.00	35.23	C
ATOM	1413	O	GLY A 190	8.748	20.857	29.930	1.00	35.26	O

ATOM	1414	N	MET	A	191	8.458	22.208	31.696	1.00	35.18	N
ATOM	1415	CA	MET	A	191	7.063	22.463	31.447	1.00	35.16	C
ATOM	1416	C	MET	A	191	6.916	23.584	30.446	1.00	35.00	C
ATOM	1417	O	MET	A	191	7.795	24.426	30.316	1.00	35.17	O
ATOM	1418	CB	MET	A	191	6.379	22.849	32.750	1.00	35.13	C
ATOM	1419	CG	MET	A	191	6.277	21.691	33.710	1.00	35.63	C
ATOM	1420	SD	MET	A	191	5.756	22.150	35.360	1.00	35.65	S
ATOM	1421	CE	MET	A	191	4.222	22.827	35.017	1.00	36.16	C
ATOM	1422	N	GLU	A	192	5.804	23.573	29.727	1.00	34.83	N
ATOM	1423	CA	GLU	A	192	5.492	24.626	28.791	1.00	34.74	C
ATOM	1424	C	GLU	A	192	5.585	25.955	29.511	1.00	34.72	C
ATOM	1425	O	GLU	A	192	5.184	26.067	30.674	1.00	34.67	O
ATOM	1426	CB	GLU	A	192	4.085	24.447	28.264	1.00	34.75	C
ATOM	1427	CG	GLU	A	192	3.029	24.525	29.348	1.00	35.09	C
ATOM	1428	CD	GLU	A	192	1.669	24.117	28.842	1.00	34.97	C
ATOM	1429	OE1	GLU	A	192	1.609	23.503	27.756	1.00	33.58	O
ATOM	1430	OE2	GLU	A	192	0.672	24.422	29.530	1.00	35.72	O
ATOM	1431	N	GLY	A	193	6.121	26.953	28.818	1.00	34.53	N
ATOM	1432	CA	GLY	A	193	6.265	28.279	29.378	1.00	34.63	C
ATOM	1433	C	GLY	A	193	7.528	28.542	30.180	1.00	34.49	C
ATOM	1434	O	GLY	A	193	7.864	29.694	30.434	1.00	34.57	O
ATOM	1435	N	ASN	A	194	8.224	27.493	30.594	1.00	34.27	N
ATOM	1436	CA	ASN	A	194	9.441	27.672	31.367	1.00	34.06	C
ATOM	1437	C	ASN	A	194	10.466	28.492	30.609	1.00	33.93	C
ATOM	1438	O	ASN	A	194	10.552	28.415	29.392	1.00	33.84	O
ATOM	1439	CB	ASN	A	194	10.051	26.316	31.721	1.00	33.99	C
ATOM	1440	CG	ASN	A	194	9.314	25.625	32.836	1.00	33.19	C
ATOM	1441	OD1	ASN	A	194	8.257	26.081	33.265	1.00	33.07	O
ATOM	1442	ND2	ASN	A	194	9.869	24.528	33.323	1.00	31.93	N
ATOM	1443	N	VAL	A	195	11.243	29.285	31.332	1.00	33.86	N
ATOM	1444	CA	VAL	A	195	12.288	30.069	30.707	1.00	33.70	C
ATOM	1445	C	VAL	A	195	13.621	29.969	31.396	1.00	33.38	C
ATOM	1446	O	VAL	A	195	13.721	30.115	32.601	1.00	33.75	O
ATOM	1447	CB	VAL	A	195	11.964	31.552	30.779	1.00	33.96	C
ATOM	1448	CG1	VAL	A	195	13.151	32.389	30.253	1.00	34.32	C
ATOM	1449	CG2	VAL	A	195	10.693	31.854	30.032	1.00	34.15	C
ATOM	1450	N	THR	A	196	14.666	29.764	30.621	1.00	33.04	N
ATOM	1451	CA	THR	A	196	15.991	29.883	31.161	1.00	32.60	C
ATOM	1452	C	THR	A	196	16.438	31.256	30.723	1.00	32.21	C
ATOM	1453	O	THR	A	196	16.584	31.500	29.530	1.00	31.64	O
ATOM	1454	CB	THR	A	196	16.887	28.836	30.586	1.00	32.68	C
ATOM	1455	OG1	THR	A	196	16.466	27.543	31.048	1.00	33.09	O
ATOM	1456	CG2	THR	A	196	18.281	29.003	31.124	1.00	32.63	C
ATOM	1457	N	PRO	A	197	16.585	32.177	31.672	1.00	31.97	N
ATOM	1458	CA	PRO	A	197	17.000	33.530	31.350	1.00	31.74	C
ATOM	1459	C	PRO	A	197	18.385	33.564	30.816	1.00	31.92	C
ATOM	1460	O	PRO	A	197	19.215	32.717	31.129	1.00	32.52	O
ATOM	1461	CB	PRO	A	197	16.950	34.234	32.682	1.00	31.38	C
ATOM	1462	CG	PRO	A	197	16.056	33.508	33.418	1.00	31.84	C
ATOM	1463	CD	PRO	A	197	16.305	32.055	33.107	1.00	32.08	C
ATOM	1464	N	ALA	A	198	18.615	34.592	30.024	1.00	32.06	N
ATOM	1465	CA	ALA	A	198	19.846	34.787	29.311	1.00	32.28	C
ATOM	1466	C	ALA	A	198	21.101	34.690	30.162	1.00	32.42	C
ATOM	1467	O	ALA	A	198	21.213	35.305	31.220	1.00	32.03	O
ATOM	1468	CB	ALA	A	198	19.798	36.131	28.625	1.00	32.42	C
ATOM	1469	N	HIS	A	199	22.064	33.953	29.630	1.00	32.77	N
ATOM	1470	CA	HIS	A	199	23.339	33.737	30.271	1.00	33.31	C
ATOM	1471	C	HIS	A	199	24.308	33.252	29.220	1.00	33.76	C
ATOM	1472	O	HIS	A	199	23.902	33.013	28.084	1.00	34.10	O
ATOM	1473	CB	HIS	A	199	23.194	32.622	31.275	1.00	33.24	C
ATOM	1474	CG	HIS	A	199	22.879	31.311	30.639	1.00	32.59	C

ATOM	1475	ND1	HIS	A	199	21.602	30.958	30.269	1.00	31.40	N
ATOM	1476	CD2	HIS	A	199	23.679	30.292	30.253	1.00	32.85	C
ATOM	1477	CE1	HIS	A	199	21.624	29.761	29.714	1.00	32.37	C
ATOM	1478	NE2	HIS	A	199	22.871	29.330	29.698	1.00	32.78	N
ATOM	1479	N	TYR	A	200	25.575	33.093	29.601	1.00	34.35	N
ATOM	1480	CA	TYR	A	200	26.580	32.502	28.712	1.00	34.92	C
ATOM	1481	C	TYR	A	200	27.256	31.350	29.431	1.00	35.15	C
ATOM	1482	O	TYR	A	200	27.285	31.318	30.661	1.00	35.76	O
ATOM	1483	CB	TYR	A	200	27.599	33.510	28.175	1.00	34.90	C
ATOM	1484	CG	TYR	A	200	28.586	34.096	29.156	1.00	34.89	C
ATOM	1485	CD1	TYR	A	200	29.825	33.519	29.359	1.00	33.02	C
ATOM	1486	CD2	TYR	A	200	28.307	35.292	29.802	1.00	36.04	C
ATOM	1487	CE1	TYR	A	200	30.731	34.079	30.224	1.00	33.25	C
ATOM	1488	CE2	TYR	A	200	29.200	35.857	30.664	1.00	35.45	C
ATOM	1489	CZ	TYR	A	200	30.410	35.253	30.882	1.00	34.40	C
ATOM	1490	OH	TYR	A	200	31.301	35.855	31.739	1.00	32.62	O
ATOM	1491	N	ASP	A	201	27.747	30.387	28.654	1.00	35.04	N
ATOM	1492	CA	ASP	A	201	28.386	29.189	29.182	1.00	34.82	C
ATOM	1493	C	ASP	A	201	29.818	29.220	28.652	1.00	34.87	C
ATOM	1494	O	ASP	A	201	30.047	29.734	27.588	1.00	34.81	O
ATOM	1495	CB	ASP	A	201	27.636	27.925	28.727	1.00	34.49	C
ATOM	1496	CG	ASP	A	201	26.212	27.854	29.256	1.00	34.63	C
ATOM	1497	OD1	ASP	A	201	26.046	27.953	30.480	1.00	33.52	O
ATOM	1498	OD2	ASP	A	201	25.191	27.680	28.533	1.00	36.84	O
ATOM	1499	N	GLU	A	202	30.795	28.727	29.399	1.00	35.74	N
ATOM	1500	CA	GLU	A	202	32.176	28.699	28.899	1.00	36.38	C
ATOM	1501	C	GLU	A	202	32.528	27.441	28.126	1.00	36.30	C
ATOM	1502	O	GLU	A	202	33.679	27.019	28.126	1.00	37.04	O
ATOM	1503	CB	GLU	A	202	33.172	28.830	30.048	1.00	36.61	C
ATOM	1504	CG	GLU	A	202	33.100	30.185	30.725	1.00	37.81	C
ATOM	1505	CD	GLU	A	202	33.960	30.276	31.949	1.00	38.93	C
ATOM	1506	OE1	GLU	A	202	33.525	29.783	33.020	1.00	39.85	O
ATOM	1507	OE2	GLU	A	202	35.055	30.858	31.832	1.00	40.19	O
ATOM	1508	N	GLN	A	203	31.556	26.840	27.462	1.00	35.68	N
ATOM	1509	CA	GLN	A	203	31.815	25.643	26.718	1.00	35.24	C
ATOM	1510	C	GLN	A	203	31.107	25.754	25.406	1.00	34.57	C
ATOM	1511	O	GLN	A	203	30.218	26.586	25.256	1.00	34.62	O
ATOM	1512	CB	GLN	A	203	31.317	24.446	27.514	1.00	35.59	C
ATOM	1513	CG	GLN	A	203	32.184	24.236	28.734	1.00	37.42	C
ATOM	1514	CD	GLN	A	203	32.234	22.819	29.233	1.00	38.70	C
ATOM	1515	OE1	GLN	A	203	32.228	21.860	28.458	1.00	39.80	O
ATOM	1516	NE2	GLN	A	203	32.326	22.680	30.541	1.00	41.01	N
ATOM	1517	N	GLN	A	204	31.524	24.939	24.443	1.00	33.68	N
ATOM	1518	CA	GLN	A	204	30.873	24.899	23.165	1.00	32.97	C
ATOM	1519	C	GLN	A	204	29.714	23.959	23.373	1.00	32.67	C
ATOM	1520	O	GLN	A	204	29.838	22.982	24.082	1.00	32.33	O
ATOM	1521	CB	GLN	A	204	31.793	24.352	22.093	1.00	32.85	C
ATOM	1522	CG	GLN	A	204	33.042	25.165	21.819	1.00	32.50	C
ATOM	1523	CD	GLN	A	204	32.786	26.454	21.057	1.00	30.59	C
ATOM	1524	OE1	GLN	A	204	31.656	26.830	20.848	1.00	31.27	O
ATOM	1525	NE2	GLN	A	204	33.843	27.132	20.664	1.00	29.18	N
ATOM	1526	N	ASN	A	205	28.583	24.247	22.755	1.00	32.81	N
ATOM	1527	CA	ASN	A	205	27.393	23.446	22.980	1.00	32.85	C
ATOM	1528	C	ASN	A	205	26.594	23.059	21.733	1.00	32.80	C
ATOM	1529	O	ASN	A	205	26.147	23.921	20.968	1.00	32.72	O
ATOM	1530	CB	ASN	A	205	26.484	24.246	23.919	1.00	32.83	C
ATOM	1531	CG	ASN	A	205	25.178	23.566	24.205	1.00	32.43	C
ATOM	1532	OD1	ASN	A	205	24.914	22.465	23.732	1.00	32.78	O
ATOM	1533	ND2	ASN	A	205	24.339	24.228	24.993	1.00	31.52	N
ATOM	1534	N	PHE	A	206	26.427	21.763	21.515	1.00	32.36	N
ATOM	1535	CA	PHE	A	206	25.454	21.326	20.541	1.00	32.02	C

ATOM	1536	C	PHE	A	206	24.167	20.977	21.301	1.00	31.59	C
ATOM	1537	O	PHE	A	206	24.144	20.067	22.096	1.00	31.13	O
ATOM	1538	CB	PHE	A	206	25.972	20.159	19.754	1.00	32.17	C
ATOM	1539	CG	PHE	A	206	26.844	20.554	18.639	1.00	32.42	C
ATOM	1540	CD1	PHE	A	206	26.364	21.354	17.630	1.00	33.16	C
ATOM	1541	CD2	PHE	A	206	28.149	20.132	18.600	1.00	33.19	C
ATOM	1542	CE1	PHE	A	206	27.174	21.721	16.600	1.00	33.34	C
ATOM	1543	CE2	PHE	A	206	28.963	20.487	17.580	1.00	32.88	C
ATOM	1544	CZ	PHE	A	206	28.485	21.279	16.574	1.00	33.50	C
ATOM	1545	N	PHE	A	207	23.104	21.717	21.012	1.00	31.65	N
ATOM	1546	CA	PHE	A	207	21.829	21.671	21.726	1.00	31.71	C
ATOM	1547	C	PHE	A	207	20.822	20.825	20.924	1.00	31.35	C
ATOM	1548	O	PHE	A	207	20.289	21.292	19.930	1.00	30.81	O
ATOM	1549	CB	PHE	A	207	21.391	23.148	21.883	1.00	31.56	C
ATOM	1550	CG	PHE	A	207	20.118	23.404	22.683	1.00	31.43	C
ATOM	1551	CD1	PHE	A	207	18.926	23.676	22.036	1.00	32.30	C
ATOM	1552	CD2	PHE	A	207	20.150	23.519	24.058	1.00	31.29	C
ATOM	1553	CE1	PHE	A	207	17.781	23.982	22.748	1.00	32.37	C
ATOM	1554	CE2	PHE	A	207	19.006	23.832	24.776	1.00	31.14	C
ATOM	1555	CZ	PHE	A	207	17.828	24.062	24.122	1.00	32.27	C
ATOM	1556	N	ALA	A	208	20.554	19.593	21.368	1.00	31.31	N
ATOM	1557	CA	ALA	A	208	19.685	18.672	20.611	1.00	31.33	C
ATOM	1558	C	ALA	A	208	18.264	18.543	21.138	1.00	31.59	C
ATOM	1559	O	ALA	A	208	18.016	17.860	22.133	1.00	31.13	O
ATOM	1560	CB	ALA	A	208	20.293	17.334	20.537	1.00	30.98	C
ATOM	1561	N	GLN	A	209	17.340	19.169	20.409	1.00	32.00	N
ATOM	1562	CA	GLN	A	209	15.935	19.183	20.751	1.00	32.37	C
ATOM	1563	C	GLN	A	209	15.288	17.867	20.358	1.00	32.76	C
ATOM	1564	O	GLN	A	209	15.492	17.360	19.247	1.00	32.19	O
ATOM	1565	CB	GLN	A	209	15.245	20.332	20.031	1.00	32.55	C
ATOM	1566	CG	GLN	A	209	13.802	20.589	20.470	1.00	32.44	C
ATOM	1567	CD	GLN	A	209	13.689	20.978	21.925	1.00	32.27	C
ATOM	1568	OE1	GLN	A	209	14.699	21.254	22.587	1.00	31.73	O
ATOM	1569	NE2	GLN	A	209	12.457	21.000	22.436	1.00	32.63	N
ATOM	1570	N	ILE	A	210	14.467	17.360	21.272	1.00	33.20	N
ATOM	1571	CA	ILE	A	210	13.907	16.030	21.165	1.00	33.55	C
ATOM	1572	C	ILE	A	210	12.399	15.998	21.195	1.00	33.37	C
ATOM	1573	O	ILE	A	210	11.788	15.337	20.369	1.00	33.25	O
ATOM	1574	CB	ILE	A	210	14.456	15.210	22.302	1.00	33.79	C
ATOM	1575	CG1	ILE	A	210	15.900	14.837	21.980	1.00	34.71	C
ATOM	1576	CG2	ILE	A	210	13.604	13.982	22.528	1.00	33.99	C
ATOM	1577	CD1	ILE	A	210	16.679	14.375	23.191	1.00	35.48	C
ATOM	1578	N	LYS	A	211	11.804	16.688	22.156	1.00	33.27	N
ATOM	1579	CA	LYS	A	211	10.365	16.739	22.263	1.00	33.48	C
ATOM	1580	C	LYS	A	211	9.960	18.146	22.559	1.00	33.68	C
ATOM	1581	O	LYS	A	211	10.545	18.789	23.407	1.00	34.10	O
ATOM	1582	CB	LYS	A	211	9.867	15.875	23.405	1.00	33.57	C
ATOM	1583	CG	LYS	A	211	8.355	15.708	23.409	1.00	34.06	C
ATOM	1584	CD	LYS	A	211	7.840	15.250	24.752	1.00	34.79	C
ATOM	1585	CE	LYS	A	211	6.635	14.324	24.635	1.00	35.65	C
ATOM	1586	NZ	LYS	A	211	5.876	14.398	23.369	1.00	36.67	N
ATOM	1587	N	GLY	A	212	8.938	18.626	21.876	1.00	34.06	N
ATOM	1588	CA	GLY	A	212	8.460	19.969	22.103	1.00	34.09	C
ATOM	1589	C	GLY	A	212	9.270	20.966	21.314	1.00	34.27	C
ATOM	1590	O	GLY	A	212	10.156	20.599	20.532	1.00	34.27	O
ATOM	1591	N	TYR	A	213	8.955	22.237	21.523	1.00	34.23	N
ATOM	1592	CA	TYR	A	213	9.640	23.311	20.843	1.00	34.12	C
ATOM	1593	C	TYR	A	213	10.216	24.318	21.824	1.00	33.81	C
ATOM	1594	O	TYR	A	213	9.602	24.652	22.835	1.00	33.09	O
ATOM	1595	CB	TYR	A	213	8.665	23.994	19.899	1.00	34.42	C
ATOM	1596	CG	TYR	A	213	8.257	23.091	18.789	1.00	35.40	C

ATOM	1597	CD1	TYR	A	213	7.228	22.163	18.955	1.00	36.80	C
ATOM	1598	CD2	TYR	A	213	8.929	23.122	17.586	1.00	36.00	C
ATOM	1599	CE1	TYR	A	213	6.873	21.323	17.944	1.00	36.82	C
ATOM	1600	CE2	TYR	A	213	8.579	22.292	16.567	1.00	37.06	C
ATOM	1601	CZ	TYR	A	213	7.560	21.390	16.743	1.00	37.82	C
ATOM	1602	OH	TYR	A	213	7.242	20.568	15.688	1.00	40.76	O
ATOM	1603	N	LYS	A	214	11.410	24.804	21.517	1.00	33.91	N
ATOM	1604	CA	LYS	A	214	12.046	25.809	22.359	1.00	33.87	C
ATOM	1605	C	LYS	A	214	12.479	27.033	21.551	1.00	33.71	C
ATOM	1606	O	LYS	A	214	13.173	26.918	20.538	1.00	33.58	O
ATOM	1607	CB	LYS	A	214	13.237	25.208	23.101	1.00	33.64	C
ATOM	1608	CG	LYS	A	214	12.881	24.443	24.364	1.00	33.56	C
ATOM	1609	CD	LYS	A	214	14.126	23.861	24.973	1.00	33.29	C
ATOM	1610	CE	LYS	A	214	14.001	23.615	26.458	1.00	33.54	C
ATOM	1611	NZ	LYS	A	214	15.346	23.419	27.130	1.00	32.27	N
ATOM	1612	N	ARG	A	215	12.043	28.205	21.986	1.00	33.56	N
ATOM	1613	CA	ARG	A	215	12.479	29.427	21.339	1.00	33.91	C
ATOM	1614	C	ARG	A	215	13.816	29.800	21.944	1.00	33.54	C
ATOM	1615	O	ARG	A	215	13.946	29.923	23.146	1.00	32.78	O
ATOM	1616	CB	ARG	A	215	11.482	30.552	21.558	1.00	34.15	C
ATOM	1617	CG	ARG	A	215	11.865	31.844	20.885	1.00	34.51	C
ATOM	1618	CD	ARG	A	215	11.287	33.028	21.593	1.00	34.99	C
ATOM	1619	NE	ARG	A	215	11.381	34.255	20.823	1.00	35.43	N
ATOM	1620	CZ	ARG	A	215	10.688	35.346	21.103	1.00	35.27	C
ATOM	1621	NH1	ARG	A	215	9.860	35.371	22.144	1.00	34.68	N
ATOM	1622	NH2	ARG	A	215	10.829	36.415	20.344	1.00	35.16	N
ATOM	1623	N	CYS	A	216	14.810	29.962	21.092	1.00	33.76	N
ATOM	1624	CA	CYS	A	216	16.152	30.238	21.542	1.00	33.91	C
ATOM	1625	C	CYS	A	216	16.555	31.613	21.068	1.00	34.31	C
ATOM	1626	O	CYS	A	216	16.534	31.885	19.872	1.00	34.69	O
ATOM	1627	CB	CYS	A	216	17.099	29.189	20.968	1.00	33.84	C
ATOM	1628	SG	CYS	A	216	16.655	27.490	21.397	1.00	32.85	S
ATOM	1629	N	ILE	A	217	16.886	32.494	22.004	1.00	34.57	N
ATOM	1630	CA	ILE	A	217	17.335	33.830	21.648	1.00	34.82	C
ATOM	1631	C	ILE	A	217	18.785	33.999	22.046	1.00	34.50	C
ATOM	1632	O	ILE	A	217	19.136	33.839	23.213	1.00	34.30	O
ATOM	1633	CB	ILE	A	217	16.475	34.890	22.324	1.00	35.07	C
ATOM	1634	CG1	ILE	A	217	15.003	34.652	22.001	1.00	35.20	C
ATOM	1635	CG2	ILE	A	217	16.881	36.259	21.844	1.00	35.38	C
ATOM	1636	CD1	ILE	A	217	14.086	35.585	22.722	1.00	36.16	C
ATOM	1637	N	LEU	A	218	19.620	34.315	21.060	1.00	34.32	N
ATOM	1638	CA	LEU	A	218	21.052	34.443	21.277	1.00	34.44	C
ATOM	1639	C	LEU	A	218	21.565	35.844	21.017	1.00	34.27	C
ATOM	1640	O	LEU	A	218	21.074	36.561	20.148	1.00	34.59	O
ATOM	1641	CB	LEU	A	218	21.818	33.476	20.365	1.00	34.57	C
ATOM	1642	CG	LEU	A	218	21.953	32.036	20.844	1.00	34.62	C
ATOM	1643	CD1	LEU	A	218	20.614	31.435	21.124	1.00	34.96	C
ATOM	1644	CD2	LEU	A	218	22.638	31.235	19.789	1.00	35.31	C
ATOM	1645	N	PHE	A	219	22.594	36.200	21.768	1.00	33.79	N
ATOM	1646	CA	PHE	A	219	23.258	37.462	21.632	1.00	33.40	C
ATOM	1647	C	PHE	A	219	24.730	37.155	21.617	1.00	33.29	C
ATOM	1648	O	PHE	A	219	25.222	36.434	22.463	1.00	33.45	O
ATOM	1649	CB	PHE	A	219	22.964	38.330	22.832	1.00	33.37	C
ATOM	1650	CG	PHE	A	219	21.509	38.586	23.051	1.00	33.91	C
ATOM	1651	CD1	PHE	A	219	20.852	39.607	22.379	1.00	33.73	C
ATOM	1652	CD2	PHE	A	219	20.794	37.812	23.942	1.00	33.70	C
ATOM	1653	CE1	PHE	A	219	19.512	39.832	22.600	1.00	33.52	C
ATOM	1654	CE2	PHE	A	219	19.460	38.038	24.156	1.00	33.47	C
ATOM	1655	CZ	PHE	A	219	18.818	39.045	23.486	1.00	33.23	C
ATOM	1656	N	PRO	A	220	25.438	37.683	20.642	1.00	33.46	N
ATOM	1657	CA	PRO	A	220	26.888	37.496	20.536	1.00	33.50	C

ATOM	1658	C	PRO	A	220	27.675	38.131	21.699	1.00	33.51	C
ATOM	1659	O	PRO	A	220	27.185	39.049	22.363	1.00	33.18	O
ATOM	1660	CB	PRO	A	220	27.246	38.169	19.211	1.00	33.51	C
ATOM	1661	CG	PRO	A	220	25.977	38.683	18.629	1.00	33.41	C
ATOM	1662	CD	PRO	A	220	24.869	38.462	19.544	1.00	33.33	C
ATOM	1663	N	PRO	A	221	28.886	37.627	21.942	1.00	33.38	N
ATOM	1664	CA	PRO	A	221	29.731	38.102	23.044	1.00	33.30	C
ATOM	1665	C	PRO	A	221	29.966	39.581	23.081	1.00	33.55	C
ATOM	1666	O	PRO	A	221	30.043	40.151	24.147	1.00	33.66	O
ATOM	1667	CB	PRO	A	221	31.033	37.375	22.799	1.00	33.11	C
ATOM	1668	CG	PRO	A	221	30.589	36.128	22.147	1.00	33.06	C
ATOM	1669	CD	PRO	A	221	29.523	36.522	21.207	1.00	32.91	C
ATOM	1670	N	ASP	A	222	30.009	40.217	21.933	1.00	34.30	N
ATOM	1671	CA	ASP	A	222	30.277	41.628	21.904	1.00	34.71	C
ATOM	1672	C	ASP	A	222	29.073	42.419	22.382	1.00	34.56	C
ATOM	1673	O	ASP	A	222	29.048	43.634	22.241	1.00	34.68	O
ATOM	1674	CB	ASP	A	222	30.679	42.055	20.502	1.00	34.79	C
ATOM	1675	CG	ASP	A	222	29.508	42.228	19.608	1.00	36.26	C
ATOM	1676	OD1	ASP	A	222	28.387	41.897	20.028	1.00	38.21	O
ATOM	1677	OD2	ASP	A	222	29.601	42.692	18.462	1.00	40.87	O
ATOM	1678	N	GLN	A	223	28.062	41.760	22.930	1.00	34.48	N
ATOM	1679	CA	GLN	A	223	26.965	42.532	23.497	1.00	34.78	C
ATOM	1680	C	GLN	A	223	27.031	42.541	25.011	1.00	34.33	C
ATOM	1681	O	GLN	A	223	26.077	42.878	25.686	1.00	34.70	O
ATOM	1682	CB	GLN	A	223	25.595	42.133	22.940	1.00	34.87	C
ATOM	1683	CG	GLN	A	223	25.364	42.825	21.599	1.00	36.55	C
ATOM	1684	CD	GLN	A	223	23.990	42.635	21.020	1.00	39.56	C
ATOM	1685	OE1	GLN	A	223	22.986	42.824	21.701	1.00	42.40	O
ATOM	1686	NE2	GLN	A	223	23.936	42.294	19.742	1.00	41.89	N
ATOM	1687	N	PHE	A	224	28.198	42.219	25.534	1.00	34.13	N
ATOM	1688	CA	PHE	A	224	28.437	42.270	26.965	1.00	34.17	C
ATOM	1689	C	PHE	A	224	27.941	43.571	27.570	1.00	34.88	C
ATOM	1690	O	PHE	A	224	27.310	43.549	28.622	1.00	35.50	O
ATOM	1691	CB	PHE	A	224	29.932	42.159	27.224	1.00	33.66	C
ATOM	1692	CG	PHE	A	224	30.305	42.027	28.661	1.00	32.69	C
ATOM	1693	CD1	PHE	A	224	30.429	43.137	29.476	1.00	33.11	C
ATOM	1694	CD2	PHE	A	224	30.609	40.790	29.191	1.00	31.85	C
ATOM	1695	CE1	PHE	A	224	30.821	43.003	30.817	1.00	31.76	C
ATOM	1696	CE2	PHE	A	224	30.993	40.661	30.498	1.00	31.29	C
ATOM	1697	CZ	PHE	A	224	31.098	41.775	31.316	1.00	30.73	C
ATOM	1698	N	GLU	A	225	28.235	44.706	26.930	1.00	35.46	N
ATOM	1699	CA	GLU	A	225	27.852	45.994	27.492	1.00	35.85	C
ATOM	1700	C	GLU	A	225	26.362	46.133	27.608	1.00	35.42	C
ATOM	1701	O	GLU	A	225	25.873	46.955	28.386	1.00	35.16	O
ATOM	1702	CB	GLU	A	225	28.401	47.175	26.688	1.00	36.59	C
ATOM	1703	CG	GLU	A	225	29.892	47.377	26.917	1.00	39.78	C
ATOM	1704	CD	GLU	A	225	30.357	48.822	27.125	1.00	43.65	C
ATOM	1705	OE1	GLU	A	225	29.937	49.524	28.099	1.00	44.64	O
ATOM	1706	OE2	GLU	A	225	31.224	49.234	26.319	1.00	47.74	O
ATOM	1707	N	CYS	A	226	25.633	45.329	26.851	1.00	35.07	N
ATOM	1708	CA	CYS	A	226	24.192	45.453	26.842	1.00	34.93	C
ATOM	1709	C	CYS	A	226	23.473	44.498	27.770	1.00	34.91	C
ATOM	1710	O	CYS	A	226	22.266	44.654	28.019	1.00	34.86	O
ATOM	1711	CB	CYS	A	226	23.681	45.196	25.448	1.00	34.80	C
ATOM	1712	SG	CYS	A	226	24.135	46.461	24.291	1.00	34.76	S
ATOM	1713	N	LEU	A	227	24.191	43.521	28.296	1.00	34.47	N
ATOM	1714	CA	LEU	A	227	23.509	42.483	29.025	1.00	34.57	C
ATOM	1715	C	LEU	A	227	23.815	42.408	30.503	1.00	34.08	C
ATOM	1716	O	LEU	A	227	23.122	41.725	31.235	1.00	33.57	O
ATOM	1717	CB	LEU	A	227	23.739	41.162	28.313	1.00	34.97	C
ATOM	1718	CG	LEU	A	227	22.883	41.127	27.047	1.00	36.04	C

ATOM	1780	CB	HIS	A	234	27.221	37.276	42.024	1.00	33.15	C
ATOM	1781	CG	HIS	A	234	25.817	37.383	42.531	1.00	32.96	C
ATOM	1782	ND1	HIS	A	234	24.767	36.704	41.951	1.00	32.53	N
ATOM	1783	CD2	HIS	A	234	25.268	38.191	43.471	1.00	32.12	C
ATOM	1784	CE1	HIS	A	234	23.639	37.056	42.540	1.00	32.45	C
ATOM	1785	NE2	HIS	A	234	23.916	37.953	43.470	1.00	32.09	N
ATOM	1786	N	PRO	A	235	26.571	34.263	42.827	1.00	33.15	N
ATOM	1787	CA	PRO	A	235	25.985	32.996	42.410	1.00	32.79	C
ATOM	1788	C	PRO	A	235	25.386	33.045	41.026	1.00	32.62	C
ATOM	1789	O	PRO	A	235	25.210	31.990	40.440	1.00	32.65	O
ATOM	1790	CB	PRO	A	235	24.897	32.773	43.450	1.00	33.00	C
ATOM	1791	CG	PRO	A	235	25.412	33.500	44.672	1.00	32.79	C
ATOM	1792	CD	PRO	A	235	26.049	34.725	44.128	1.00	33.00	C
ATOM	1793	N	CYS	A	236	25.095	34.226	40.493	1.00	32.39	N
ATOM	1794	CA	CYS	A	236	24.487	34.286	39.185	1.00	32.11	C
ATOM	1795	C	CYS	A	236	25.529	34.640	38.139	1.00	32.34	C
ATOM	1796	O	CYS	A	236	25.217	35.034	37.018	1.00	32.23	O
ATOM	1797	CB	CYS	A	236	23.270	35.192	39.206	1.00	32.02	C
ATOM	1798	SG	CYS	A	236	21.990	34.545	40.326	1.00	32.09	S
ATOM	1799	N	ASP	A	237	26.789	34.456	38.511	1.00	32.71	N
ATOM	1800	CA	ASP	A	237	27.883	34.609	37.576	1.00	33.21	C
ATOM	1801	C	ASP	A	237	27.518	34.022	36.211	1.00	33.63	C
ATOM	1802	O	ASP	A	237	27.042	32.911	36.095	1.00	33.33	O
ATOM	1803	CB	ASP	A	237	29.139	33.964	38.132	1.00	33.01	C
ATOM	1804	CG	ASP	A	237	30.283	33.981	37.151	1.00	33.76	C
ATOM	1805	OD1	ASP	A	237	30.340	34.874	36.279	1.00	34.52	O
ATOM	1806	OD2	ASP	A	237	31.193	33.135	37.181	1.00	36.09	O
ATOM	1807	N	ARG	A	238	27.731	34.821	35.180	1.00	34.68	N
ATOM	1808	CA	ARG	A	238	27.414	34.451	33.810	1.00	35.15	C
ATOM	1809	C	ARG	A	238	25.962	34.690	33.375	1.00	34.98	C
ATOM	1810	O	ARG	A	238	25.664	34.543	32.197	1.00	34.26	O
ATOM	1811	CB	ARG	A	238	27.819	33.020	33.561	1.00	35.64	C
ATOM	1812	CG	ARG	A	238	29.286	32.847	33.620	1.00	36.86	C
ATOM	1813	CD	ARG	A	238	29.682	31.474	33.268	1.00	38.59	C
ATOM	1814	NE	ARG	A	238	29.321	30.549	34.338	1.00	41.64	N
ATOM	1815	CZ	ARG	A	238	28.265	29.755	34.286	1.00	43.17	C
ATOM	1816	NH1	ARG	A	238	27.459	29.802	33.218	1.00	44.38	N
ATOM	1817	NH2	ARG	A	238	28.006	28.923	35.291	1.00	42.18	N
ATOM	1818	N	GLN	A	239	25.067	35.044	34.303	1.00	35.13	N
ATOM	1819	CA	GLN	A	239	23.683	35.360	33.920	1.00	35.13	C
ATOM	1820	C	GLN	A	239	23.558	36.872	33.825	1.00	34.56	C
ATOM	1821	O	GLN	A	239	24.240	37.587	34.539	1.00	34.65	O
ATOM	1822	CB	GLN	A	239	22.646	34.839	34.921	1.00	35.27	C
ATOM	1823	CG	GLN	A	239	22.952	33.510	35.566	1.00	37.30	C
ATOM	1824	CD	GLN	A	239	23.270	32.431	34.552	1.00	41.59	C
ATOM	1825	OE1	GLN	A	239	22.418	32.072	33.726	1.00	45.42	O
ATOM	1826	NE2	GLN	A	239	24.491	31.899	34.609	1.00	42.92	N
ATOM	1827	N	SER	A	240	22.705	37.358	32.930	1.00	33.99	N
ATOM	1828	CA	SER	A	240	22.455	38.785	32.802	1.00	33.14	C
ATOM	1829	C	SER	A	240	21.613	39.254	33.957	1.00	32.53	C
ATOM	1830	O	SER	A	240	20.773	38.528	34.446	1.00	31.77	O
ATOM	1831	CB	SER	A	240	21.663	39.077	31.542	1.00	33.15	C
ATOM	1832	OG	SER	A	240	20.971	40.309	31.668	1.00	33.28	O
ATOM	1833	N	GLN	A	241	21.805	40.492	34.371	1.00	32.52	N
ATOM	1834	CA	GLN	A	241	21.022	41.025	35.476	1.00	32.54	C
ATOM	1835	C	GLN	A	241	19.711	41.629	35.001	1.00	32.44	C
ATOM	1836	O	GLN	A	241	18.872	42.019	35.804	1.00	32.16	O
ATOM	1837	CB	GLN	A	241	21.791	42.116	36.196	1.00	32.36	C
ATOM	1838	CG	GLN	A	241	22.995	41.669	36.917	1.00	32.74	C
ATOM	1839	CD	GLN	A	241	23.760	42.837	37.450	1.00	33.65	C
ATOM	1840	OE1	GLN	A	241	23.353	43.442	38.428	1.00	34.26	O

ATOM	1841	NE2	GLN	A	241	24.858	43.182	36.794	1.00	35.62	N
ATOM	1842	N	VAL	A	242	19.513	41.708	33.705	1.00	32.07	N
ATOM	1843	CA	VAL	A	242	18.357	42.412	33.255	1.00	32.29	C
ATOM	1844	C	VAL	A	242	17.162	41.521	33.265	1.00	31.93	C
ATOM	1845	O	VAL	A	242	17.221	40.442	32.734	1.00	32.93	O
ATOM	1846	CB	VAL	A	242	18.516	42.848	31.808	1.00	32.63	C
ATOM	1847	CG1	VAL	A	242	17.252	43.532	31.320	1.00	32.50	C
ATOM	1848	CG2	VAL	A	242	19.717	43.714	31.655	1.00	33.06	C
ATOM	1849	N	ASP	A	243	16.067	41.977	33.838	1.00	31.76	N
ATOM	1850	CA	ASP	A	243	14.812	41.271	33.714	1.00	31.73	C
ATOM	1851	C	ASP	A	243	14.177	41.559	32.346	1.00	31.87	C
ATOM	1852	O	ASP	A	243	13.536	42.595	32.150	1.00	31.46	O
ATOM	1853	CB	ASP	A	243	13.861	41.689	34.830	1.00	31.61	C
ATOM	1854	CG	ASP	A	243	12.488	41.049	34.708	1.00	32.08	C
ATOM	1855	OD1	ASP	A	243	12.164	40.448	33.655	1.00	30.69	O
ATOM	1856	OD2	ASP	A	243	11.654	41.103	35.635	1.00	33.90	O
ATOM	1857	N	PHE	A	244	14.326	40.623	31.410	1.00	32.16	N
ATOM	1858	CA	PHE	A	244	13.746	40.787	30.075	1.00	32.29	C
ATOM	1859	C	PHE	A	244	12.252	41.087	30.092	1.00	32.74	C
ATOM	1860	O	PHE	A	244	11.741	41.694	29.162	1.00	32.51	O
ATOM	1861	CB	PHE	A	244	13.963	39.559	29.220	1.00	31.99	C
ATOM	1862	CG	PHE	A	244	15.327	39.446	28.650	1.00	31.29	C
ATOM	1863	CD1	PHE	A	244	16.436	39.886	29.342	1.00	31.05	C
ATOM	1864	CD2	PHE	A	244	15.500	38.861	27.421	1.00	31.04	C
ATOM	1865	CE1	PHE	A	244	17.684	39.733	28.815	1.00	31.37	C
ATOM	1866	CE2	PHE	A	244	16.742	38.709	26.888	1.00	31.39	C
ATOM	1867	CZ	PHE	A	244	17.839	39.141	27.584	1.00	31.77	C
ATOM	1868	N	ASP	A	245	11.547	40.646	31.124	1.00	33.52	N
ATOM	1869	CA	ASP	A	245	10.130	40.946	31.214	1.00	34.33	C
ATOM	1870	C	ASP	A	245	9.831	42.375	31.606	1.00	34.52	C
ATOM	1871	O	ASP	A	245	8.789	42.902	31.251	1.00	34.40	O
ATOM	1872	CB	ASP	A	245	9.448	40.015	32.188	1.00	34.59	C
ATOM	1873	CG	ASP	A	245	9.361	38.639	31.658	1.00	35.47	C
ATOM	1874	OD1	ASP	A	245	9.282	38.519	30.420	1.00	36.40	O
ATOM	1875	OD2	ASP	A	245	9.384	37.624	32.383	1.00	37.65	O
ATOM	1876	N	ASN	A	246	10.737	43.000	32.339	1.00	35.03	N
ATOM	1877	CA	ASN	A	246	10.531	44.364	32.776	1.00	35.38	C
ATOM	1878	C	ASN	A	246	11.873	45.003	32.896	1.00	34.97	C
ATOM	1879	O	ASN	A	246	12.370	45.160	33.994	1.00	35.01	O
ATOM	1880	CB	ASN	A	246	9.843	44.397	34.136	1.00	35.69	C
ATOM	1881	CG	ASN	A	246	9.395	45.800	34.533	1.00	37.41	C
ATOM	1882	OD1	ASN	A	246	9.241	46.691	33.682	1.00	38.99	O
ATOM	1883	ND2	ASN	A	246	9.186	46.006	35.836	1.00	38.90	N
ATOM	1884	N	PRO	A	247	12.468	45.357	31.766	1.00	34.87	N
ATOM	1885	CA	PRO	A	247	13.816	45.931	31.766	1.00	34.87	C
ATOM	1886	C	PRO	A	247	13.868	47.295	32.407	1.00	34.83	C
ATOM	1887	O	PRO	A	247	13.038	48.159	32.127	1.00	34.82	O
ATOM	1888	CB	PRO	A	247	14.184	46.046	30.288	1.00	34.75	C
ATOM	1889	CG	PRO	A	247	13.028	45.475	29.507	1.00	34.95	C
ATOM	1890	CD	PRO	A	247	11.892	45.241	30.419	1.00	34.75	C
ATOM	1891	N	ASP	A	248	14.861	47.448	33.270	1.00	34.67	N
ATOM	1892	CA	ASP	A	248	15.112	48.654	34.003	1.00	34.76	C
ATOM	1893	C	ASP	A	248	16.205	49.419	33.294	1.00	34.89	C
ATOM	1894	O	ASP	A	248	17.395	49.240	33.568	1.00	34.59	O
ATOM	1895	CB	ASP	A	248	15.583	48.265	35.387	1.00	34.87	C
ATOM	1896	CG	ASP	A	248	15.703	49.433	36.329	1.00	35.10	C
ATOM	1897	OD1	ASP	A	248	15.958	50.583	35.902	1.00	34.24	O
ATOM	1898	OD2	ASP	A	248	15.574	49.259	37.550	1.00	36.72	O
ATOM	1899	N	TYR	A	249	15.793	50.302	32.399	1.00	35.08	N
ATOM	1900	CA	TYR	A	249	16.743	51.063	31.615	1.00	35.24	C
ATOM	1901	C	TYR	A	249	17.578	52.071	32.400	1.00	35.48	C

ATOM	1902	O	TYR	A	249	18.570	52.567	31.880	1.00	35.44	O
ATOM	1903	CB	TYR	A	249	16.021	51.741	30.465	1.00	35.05	C
ATOM	1904	CG	TYR	A	249	15.304	50.761	29.586	1.00	34.41	C
ATOM	1905	CD1	TYR	A	249	15.977	49.718	28.971	1.00	34.17	C
ATOM	1906	CD2	TYR	A	249	13.955	50.865	29.383	1.00	34.14	C
ATOM	1907	CE1	TYR	A	249	15.312	48.828	28.172	1.00	33.95	C
ATOM	1908	CE2	TYR	A	249	13.287	49.983	28.595	1.00	33.71	C
ATOM	1909	CZ	TYR	A	249	13.954	48.974	27.989	1.00	34.19	C
ATOM	1910	OH	TYR	A	249	13.232	48.113	27.194	1.00	35.64	O
ATOM	1911	N	GLU	A	250	17.207	52.393	33.631	1.00	35.82	N
ATOM	1912	CA	GLU	A	250	18.072	53.273	34.399	1.00	36.69	C
ATOM	1913	C	GLU	A	250	19.331	52.562	34.834	1.00	36.69	C
ATOM	1914	O	GLU	A	250	20.424	53.102	34.715	1.00	36.84	O
ATOM	1915	CB	GLU	A	250	17.369	53.836	35.607	1.00	37.10	C
ATOM	1916	CG	GLU	A	250	16.173	54.653	35.195	1.00	39.48	C
ATOM	1917	CD	GLU	A	250	15.559	55.362	36.359	1.00	42.67	C
ATOM	1918	OE1	GLU	A	250	16.128	55.258	37.469	1.00	45.95	O
ATOM	1919	OE2	GLU	A	250	14.529	56.030	36.160	1.00	44.93	O
ATOM	1920	N	ARG	A	251	19.202	51.344	35.332	1.00	36.55	N
ATOM	1921	CA	ARG	A	251	20.389	50.666	35.773	1.00	36.36	C
ATOM	1922	C	ARG	A	251	21.124	50.162	34.584	1.00	35.91	C
ATOM	1923	O	ARG	A	251	22.347	50.117	34.588	1.00	36.89	O
ATOM	1924	CB	ARG	A	251	20.062	49.495	36.697	1.00	36.66	C
ATOM	1925	CG	ARG	A	251	19.329	49.936	37.981	1.00	38.10	C
ATOM	1926	CD	ARG	A	251	18.848	48.824	38.916	1.00	39.00	C
ATOM	1927	NE	ARG	A	251	19.969	48.005	39.371	1.00	40.55	N
ATOM	1928	CZ	ARG	A	251	19.895	46.712	39.648	1.00	41.33	C
ATOM	1929	NH1	ARG	A	251	18.742	46.068	39.542	1.00	42.58	N
ATOM	1930	NH2	ARG	A	251	20.978	46.058	40.025	1.00	41.49	N
ATOM	1931	N	PHE	A	252	20.388	49.807	33.545	1.00	35.06	N
ATOM	1932	CA	PHE	A	252	20.988	49.065	32.455	1.00	34.37	C
ATOM	1933	C	PHE	A	252	20.653	49.675	31.138	1.00	33.82	C
ATOM	1934	O	PHE	A	252	20.048	49.042	30.278	1.00	33.57	O
ATOM	1935	CB	PHE	A	252	20.429	47.662	32.458	1.00	34.19	C
ATOM	1936	CG	PHE	A	252	20.404	47.019	33.803	1.00	33.98	C
ATOM	1937	CD1	PHE	A	252	21.559	46.855	34.532	1.00	33.98	C
ATOM	1938	CD2	PHE	A	252	19.220	46.560	34.337	1.00	32.84	C
ATOM	1939	CE1	PHE	A	252	21.519	46.238	35.760	1.00	33.51	C
ATOM	1940	CE2	PHE	A	252	19.189	45.953	35.560	1.00	32.14	C
ATOM	1941	CZ	PHE	A	252	20.327	45.789	36.269	1.00	31.96	C
ATOM	1942	N	PRO	A	253	21.087	50.903	30.959	1.00	33.27	N
ATOM	1943	CA	PRO	A	253	20.677	51.675	29.795	1.00	32.92	C
ATOM	1944	C	PRO	A	253	21.008	50.995	28.470	1.00	32.86	C
ATOM	1945	O	PRO	A	253	20.215	51.136	27.541	1.00	32.58	O
ATOM	1946	CB	PRO	A	253	21.464	52.957	29.962	1.00	32.58	C
ATOM	1947	CG	PRO	A	253	22.584	52.562	30.808	1.00	32.44	C
ATOM	1948	CD	PRO	A	253	22.038	51.640	31.802	1.00	32.67	C
ATOM	1949	N	ASN	A	254	22.108	50.253	28.359	1.00	32.60	N
ATOM	1950	CA	ASN	A	254	22.419	49.726	27.037	1.00	32.87	C
ATOM	1951	C	ASN	A	254	21.564	48.569	26.615	1.00	32.59	C
ATOM	1952	O	ASN	A	254	21.671	48.088	25.503	1.00	32.69	O
ATOM	1953	CB	ASN	A	254	23.892	49.414	26.852	1.00	32.72	C
ATOM	1954	CG	ASN	A	254	24.710	50.665	26.745	1.00	33.92	C
ATOM	1955	OD1	ASN	A	254	25.428	51.022	27.672	1.00	37.05	O
ATOM	1956	ND2	ASN	A	254	24.569	51.380	25.626	1.00	34.86	N
ATOM	1957	N	PHE	A	255	20.690	48.124	27.491	1.00	32.37	N
ATOM	1958	CA	PHE	A	255	19.834	47.057	27.101	1.00	32.19	C
ATOM	1959	C	PHE	A	255	18.932	47.578	26.006	1.00	32.36	C
ATOM	1960	O	PHE	A	255	18.267	46.819	25.326	1.00	32.64	O
ATOM	1961	CB	PHE	A	255	18.990	46.572	28.250	1.00	31.93	C
ATOM	1962	CG	PHE	A	255	18.249	45.354	27.918	1.00	31.43	C

ATOM	1963	CD1	PHE	A	255	18.919	44.176	27.718	1.00	33.75	C
ATOM	1964	CD2	PHE	A	255	16.909	45.392	27.718	1.00	31.23	C
ATOM	1965	CE1	PHE	A	255	18.239	43.035	27.367	1.00	34.05	C
ATOM	1966	CE2	PHE	A	255	16.230	44.275	27.380	1.00	32.00	C
ATOM	1967	CZ	PHE	A	255	16.890	43.088	27.203	1.00	32.84	C
ATOM	1968	N	GLN	A	256	18.890	48.888	25.844	1.00	32.46	N
ATOM	1969	CA	GLN	A	256	18.078	49.456	24.794	1.00	32.37	C
ATOM	1970	C	GLN	A	256	18.776	49.342	23.455	1.00	32.34	C
ATOM	1971	O	GLN	A	256	18.260	49.815	22.470	1.00	32.11	O
ATOM	1972	CB	GLN	A	256	17.777	50.924	25.061	1.00	32.28	C
ATOM	1973	CG	GLN	A	256	16.775	51.169	26.144	1.00	32.39	C
ATOM	1974	CD	GLN	A	256	16.823	52.594	26.645	1.00	32.91	C
ATOM	1975	OE1	GLN	A	256	15.830	53.303	26.588	1.00	34.66	O
ATOM	1976	NE2	GLN	A	256	17.982	53.021	27.125	1.00	32.30	N
ATOM	1977	N	ASN	A	257	19.956	48.744	23.404	1.00	32.78	N
ATOM	1978	CA	ASN	A	257	20.634	48.617	22.126	1.00	33.17	C
ATOM	1979	C	ASN	A	257	20.828	47.159	21.763	1.00	34.01	C
ATOM	1980	O	ASN	A	257	21.406	46.841	20.721	1.00	34.41	O
ATOM	1981	CB	ASN	A	257	21.998	49.310	22.140	1.00	32.88	C
ATOM	1982	CG	ASN	A	257	21.928	50.750	22.595	1.00	31.31	C
ATOM	1983	OD1	ASN	A	257	22.471	51.105	23.639	1.00	27.26	O
ATOM	1984	ND2	ASN	A	257	21.283	51.595	21.797	1.00	29.34	N
ATOM	1985	N	VAL	A	258	20.338	46.263	22.606	1.00	34.66	N
ATOM	1986	CA	VAL	A	258	20.565	44.855	22.370	1.00	35.33	C
ATOM	1987	C	VAL	A	258	19.832	44.361	21.130	1.00	35.37	C
ATOM	1988	O	VAL	A	258	18.751	44.832	20.813	1.00	34.96	O
ATOM	1989	CB	VAL	A	258	20.144	44.010	23.558	1.00	35.59	C
ATOM	1990	CG1	VAL	A	258	18.632	43.851	23.604	1.00	35.73	C
ATOM	1991	CG2	VAL	A	258	20.780	42.662	23.428	1.00	36.60	C
ATOM	1992	N	VAL	A	259	20.447	43.418	20.428	1.00	35.87	N
ATOM	1993	CA	VAL	A	259	19.853	42.844	19.230	1.00	36.40	C
ATOM	1994	C	VAL	A	259	20.125	41.355	19.148	1.00	36.66	C
ATOM	1995	O	VAL	A	259	21.282	40.943	19.102	1.00	36.47	O
ATOM	1996	CB	VAL	A	259	20.450	43.459	17.969	1.00	36.47	C
ATOM	1997	CG1	VAL	A	259	19.830	42.822	16.764	1.00	36.75	C
ATOM	1998	CG2	VAL	A	259	20.212	44.931	17.932	1.00	36.66	C
ATOM	1999	N	GLY	A	260	19.066	40.551	19.085	1.00	37.15	N
ATOM	2000	CA	GLY	A	260	19.215	39.096	19.067	1.00	37.50	C
ATOM	2001	C	GLY	A	260	19.132	38.328	17.745	1.00	37.55	C
ATOM	2002	O	GLY	A	260	18.716	38.839	16.704	1.00	37.45	O
ATOM	2003	N	TYR	A	261	19.578	37.079	17.817	1.00	37.54	N
ATOM	2004	CA	TYR	A	261	19.466	36.113	16.744	1.00	37.52	C
ATOM	2005	C	TYR	A	261	18.530	35.107	17.358	1.00	37.09	C
ATOM	2006	O	TYR	A	261	18.788	34.648	18.466	1.00	37.12	O
ATOM	2007	CB	TYR	A	261	20.796	35.429	16.481	1.00	37.79	C
ATOM	2008	CG	TYR	A	261	21.838	36.327	15.874	1.00	39.43	C
ATOM	2009	CD1	TYR	A	261	22.005	36.380	14.507	1.00	41.03	C
ATOM	2010	CD2	TYR	A	261	22.648	37.125	16.662	1.00	41.07	C
ATOM	2011	CE1	TYR	A	261	22.937	37.193	13.940	1.00	41.71	C
ATOM	2012	CE2	TYR	A	261	23.593	37.938	16.093	1.00	41.99	C
ATOM	2013	CZ	TYR	A	261	23.725	37.960	14.728	1.00	42.45	C
ATOM	2014	OH	TYR	A	261	24.650	38.762	14.121	1.00	46.29	O
ATOM	2015	N	GLU	A	262	17.430	34.773	16.699	1.00	36.56	N
ATOM	2016	CA	GLU	A	262	16.500	33.866	17.346	1.00	35.87	C
ATOM	2017	C	GLU	A	262	16.064	32.737	16.447	1.00	35.41	C
ATOM	2018	O	GLU	A	262	16.310	32.747	15.248	1.00	35.03	O
ATOM	2019	CB	GLU	A	262	15.320	34.639	17.943	1.00	35.67	C
ATOM	2020	CG	GLU	A	262	14.085	34.789	17.098	1.00	35.22	C
ATOM	2021	CD	GLU	A	262	13.039	35.601	17.821	1.00	35.62	C
ATOM	2022	OE1	GLU	A	262	13.201	36.833	17.866	1.00	38.15	O
ATOM	2023	OE2	GLU	A	262	12.074	35.031	18.367	1.00	35.12	O

ATOM	2024	N	THR	A	263	15.440	31.744	17.060	1.00	35.07	N
ATOM	2025	CA	THR	A	263	15.004	30.565	16.351	1.00	34.84	C
ATOM	2026	C	THR	A	263	14.138	29.670	17.224	1.00	34.38	C
ATOM	2027	O	THR	A	263	14.152	29.745	18.452	1.00	34.07	O
ATOM	2028	CB	THR	A	263	16.235	29.767	15.847	1.00	34.75	C
ATOM	2029	OG1	THR	A	263	15.864	28.964	14.731	1.00	35.81	O
ATOM	2030	CG2	THR	A	263	16.693	28.745	16.837	1.00	34.59	C
ATOM	2031	N	VAL	A	264	13.368	28.820	16.573	1.00	34.02	N
ATOM	2032	CA	VAL	A	264	12.597	27.854	17.306	1.00	33.95	C
ATOM	2033	C	VAL	A	264	13.054	26.460	16.918	1.00	33.60	C
ATOM	2034	O	VAL	A	264	12.957	26.065	15.762	1.00	33.27	O
ATOM	2035	CB	VAL	A	264	11.112	28.019	17.075	1.00	33.95	C
ATOM	2036	CG1	VAL	A	264	10.393	26.788	17.568	1.00	34.24	C
ATOM	2037	CG2	VAL	A	264	10.615	29.247	17.823	1.00	33.85	C
ATOM	2038	N	VAL	A	265	13.572	25.708	17.880	1.00	33.43	N
ATOM	2039	CA	VAL	A	265	13.984	24.354	17.546	1.00	33.48	C
ATOM	2040	C	VAL	A	265	12.949	23.321	17.907	1.00	32.85	C
ATOM	2041	O	VAL	A	265	12.234	23.441	18.884	1.00	32.91	O
ATOM	2042	CB	VAL	A	265	15.332	23.954	18.139	1.00	33.61	C
ATOM	2043	CG1	VAL	A	265	16.408	24.721	17.432	1.00	34.56	C
ATOM	2044	CG2	VAL	A	265	15.381	24.153	19.636	1.00	33.15	C
ATOM	2045	N	GLY	A	266	12.848	22.321	17.064	1.00	32.32	N
ATOM	2046	CA	GLY	A	266	11.938	21.240	17.331	1.00	32.27	C
ATOM	2047	C	GLY	A	266	12.634	19.905	17.310	1.00	31.78	C
ATOM	2048	O	GLY	A	266	13.852	19.799	17.124	1.00	31.15	O
ATOM	2049	N	PRO	A	267	11.827	18.872	17.454	1.00	31.56	N
ATOM	2050	CA	PRO	A	267	12.343	17.516	17.547	1.00	31.40	C
ATOM	2051	C	PRO	A	267	13.203	17.314	16.336	1.00	31.22	C
ATOM	2052	O	PRO	A	267	12.755	17.631	15.241	1.00	31.24	O
ATOM	2053	CB	PRO	A	267	11.074	16.664	17.552	1.00	30.81	C
ATOM	2054	CG	PRO	A	267	10.048	17.568	18.087	1.00	31.26	C
ATOM	2055	CD	PRO	A	267	10.358	18.900	17.451	1.00	31.85	C
ATOM	2056	N	GLY	A	268	14.445	16.891	16.536	1.00	31.06	N
ATOM	2057	CA	GLY	A	268	15.334	16.620	15.426	1.00	31.07	C
ATOM	2058	C	GLY	A	268	16.337	17.703	15.112	1.00	31.31	C
ATOM	2059	O	GLY	A	268	17.352	17.425	14.493	1.00	31.12	O
ATOM	2060	N	ASP	A	269	16.061	18.939	15.517	1.00	31.87	N
ATOM	2061	CA	ASP	A	269	16.975	20.032	15.253	1.00	32.18	C
ATOM	2062	C	ASP	A	269	18.117	20.099	16.282	1.00	32.69	C
ATOM	2063	O	ASP	A	269	17.974	19.720	17.450	1.00	32.69	O
ATOM	2064	CB	ASP	A	269	16.282	21.381	15.390	1.00	32.73	C
ATOM	2065	CG	ASP	A	269	15.094	21.583	14.478	1.00	32.07	C
ATOM	2066	OD1	ASP	A	269	15.023	21.039	13.367	1.00	33.96	O
ATOM	2067	OD2	ASP	A	269	14.191	22.368	14.806	1.00	30.81	O
ATOM	2068	N	VAL	A	270	19.234	20.667	15.854	1.00	32.97	N
ATOM	2069	CA	VAL	A	270	20.376	20.839	16.715	1.00	32.97	C
ATOM	2070	C	VAL	A	270	20.844	22.258	16.579	1.00	32.97	C
ATOM	2071	O	VAL	A	270	21.130	22.716	15.488	1.00	33.40	O
ATOM	2072	CB	VAL	A	270	21.485	19.896	16.323	1.00	32.94	C
ATOM	2073	CG1	VAL	A	270	22.755	20.207	17.066	1.00	33.16	C
ATOM	2074	CG2	VAL	A	270	21.069	18.519	16.646	1.00	33.14	C
ATOM	2075	N	LEU	A	271	20.883	22.965	17.692	1.00	32.91	N
ATOM	2076	CA	LEU	A	271	21.321	24.337	17.690	1.00	32.91	C
ATOM	2077	C	LEU	A	271	22.770	24.395	18.129	1.00	32.95	C
ATOM	2078	O	LEU	A	271	23.116	23.864	19.161	1.00	32.76	O
ATOM	2079	CB	LEU	A	271	20.468	25.159	18.656	1.00	32.64	C
ATOM	2080	CG	LEU	A	271	20.896	26.616	18.773	1.00	32.61	C
ATOM	2081	CD1	LEU	A	271	20.989	27.276	17.405	1.00	32.23	C
ATOM	2082	CD2	LEU	A	271	19.936	27.383	19.638	1.00	33.12	C
ATOM	2083	N	TYR	A	272	23.633	25.006	17.333	1.00	33.19	N
ATOM	2084	CA	TYR	A	272	24.989	25.215	17.791	1.00	33.39	C

ATOM	2085	C	TYR	A	272	25.004	26.517	18.554	1.00	33.40
ATOM	2086	O	TYR	A	272	24.834	27.560	17.950	1.00	33.42
ATOM	2087	CB	TYR	A	272	25.991	25.297	16.633	1.00	33.28
ATOM	2088	CG	TYR	A	272	27.376	25.802	17.039	1.00	32.43
ATOM	2089	CD1	TYR	A	272	28.005	25.340	18.179	1.00	31.06
ATOM	2090	CD2	TYR	A	272	28.047	26.746	16.267	1.00	32.28
ATOM	2091	CE1	TYR	A	272	29.265	25.805	18.544	1.00	30.52
ATOM	2092	CE2	TYR	A	272	29.300	27.213	16.622	1.00	30.88
ATOM	2093	CZ	TYR	A	272	29.906	26.737	17.759	1.00	29.92
ATOM	2094	OH	TYR	A	272	31.146	27.216	18.113	1.00	27.15
ATOM	2095	N	ILE	A	273	25.178	26.438	19.872	1.00	33.69
ATOM	2096	CA	ILE	A	273	25.350	27.609	20.734	1.00	33.97
ATOM	2097	C	ILE	A	273	26.830	27.759	21.068	1.00	34.43
ATOM	2098	O	ILE	A	273	27.332	27.113	21.998	1.00	34.76
ATOM	2099	CB	ILE	A	273	24.595	27.436	22.032	1.00	33.75
ATOM	2100	CG1	ILE	A	273	23.122	27.219	21.749	1.00	33.59
ATOM	2101	CG2	ILE	A	273	24.779	28.654	22.896	1.00	33.60
ATOM	2102	CD1	ILE	A	273	22.306	26.986	22.984	1.00	33.23
ATOM	2103	N	PRO	A	274	27.519	28.630	20.345	1.00	34.61
ATOM	2104	CA	PRO	A	274	28.965	28.780	20.485	1.00	34.92
ATOM	2105	C	PRO	A	274	29.348	29.378	21.803	1.00	35.17
ATOM	2106	O	PRO	A	274	28.639	30.252	22.300	1.00	35.16
ATOM	2107	CB	PRO	A	274	29.333	29.771	19.382	1.00	35.09
ATOM	2108	CG	PRO	A	274	28.113	29.929	18.551	1.00	35.02
ATOM	2109	CD	PRO	A	274	26.957	29.584	19.388	1.00	34.69
ATOM	2110	N	MET	A	275	30.476	28.927	22.339	1.00	35.69
ATOM	2111	CA	MET	A	275	30.982	29.399	23.622	1.00	36.15
ATOM	2112	C	MET	A	275	30.981	30.917	23.754	1.00	36.09
ATOM	2113	O	MET	A	275	31.316	31.635	22.805	1.00	36.23
ATOM	2114	CB	MET	A	275	32.400	28.928	23.805	1.00	36.23
ATOM	2115	CG	MET	A	275	32.813	28.928	25.229	1.00	38.02
ATOM	2116	SD	MET	A	275	34.361	28.108	25.442	1.00	42.07
ATOM	2117	CE	MET	A	275	35.330	28.680	23.895	1.00	42.34
ATOM	2118	N	TYR	A	276	30.611	31.401	24.935	1.00	35.76
ATOM	2119	CA	TYR	A	276	30.574	32.833	25.191	1.00	35.62
ATOM	2120	C	TYR	A	276	29.339	33.513	24.555	1.00	35.32
ATOM	2121	O	TYR	A	276	29.048	34.653	24.866	1.00	35.15
ATOM	2122	CB	TYR	A	276	31.921	33.510	24.804	1.00	35.75
ATOM	2123	CG	TYR	A	276	33.049	33.221	25.822	1.00	35.70
ATOM	2124	CD1	TYR	A	276	33.036	33.805	27.077	1.00	34.51
ATOM	2125	CD2	TYR	A	276	34.103	32.356	25.521	1.00	35.30
ATOM	2126	CE1	TYR	A	276	34.013	33.549	27.997	1.00	34.63

ATOM	2146	C	TRP	A	278	23.334	33.859	24.838	1.00	35.14	C
ATOM	2147	O	TRP	A	278	23.068	33.939	23.637	1.00	35.16	O
ATOM	2148	CB	TRP	A	278	24.047	36.030	25.553	1.00	34.94	C
ATOM	2149	CG	TRP	A	278	25.025	36.909	26.185	1.00	35.52	C
ATOM	2150	CD1	TRP	A	278	26.030	37.597	25.584	1.00	35.77	C
ATOM	2151	CD2	TRP	A	278	25.071	37.244	27.560	1.00	36.19	C
ATOM	2152	NE1	TRP	A	278	26.713	38.336	26.518	1.00	36.09	N
ATOM	2153	CE2	TRP	A	278	26.133	38.132	27.741	1.00	36.37	C
ATOM	2154	CE3	TRP	A	278	24.312	36.876	28.668	1.00	36.48	C
ATOM	2155	CZ2	TRP	A	278	26.463	38.637	28.976	1.00	37.12	C
ATOM	2156	CZ3	TRP	A	278	24.636	37.375	29.877	1.00	36.37	C
ATOM	2157	CH2	TRP	A	278	25.701	38.246	30.032	1.00	36.92	C
ATOM	2158	N	HIS	A	279	22.620	33.133	25.692	1.00	35.10	N
ATOM	2159	CA	HIS	A	279	21.400	32.516	25.218	1.00	35.20	C
ATOM	2160	C	HIS	A	279	20.300	32.555	26.231	1.00	34.69	C
ATOM	2161	O	HIS	A	279	20.540	32.348	27.414	1.00	34.75	O
ATOM	2162	CB	HIS	A	279	21.624	31.079	24.764	1.00	35.73	C
ATOM	2163	CG	HIS	A	279	22.403	30.236	25.727	1.00	37.15	C
ATOM	2164	ND1	HIS	A	279	23.775	30.295	25.819	1.00	39.54	N
ATOM	2165	CD2	HIS	A	279	22.010	29.283	26.609	1.00	38.10	C
ATOM	2166	CE1	HIS	A	279	24.194	29.425	26.726	1.00	39.33	C
ATOM	2167	NE2	HIS	A	279	23.143	28.797	27.221	1.00	37.68	N
ATOM	2168	N	HIS	A	280	19.092	32.795	25.721	1.00	34.17	N
ATOM	2169	CA	HIS	A	280	17.850	32.854	26.483	1.00	33.72	C
ATOM	2170	C	HIS	A	280	16.960	31.779	25.901	1.00	33.50	C
ATOM	2171	O	HIS	A	280	16.730	31.765	24.705	1.00	33.21	O
ATOM	2172	CB	HIS	A	280	17.215	34.208	26.257	1.00	33.58	C
ATOM	2173	CG	HIS	A	280	15.733	34.247	26.468	1.00	33.92	C
ATOM	2174	ND1	HIS	A	280	15.151	34.997	27.464	1.00	33.67	N
ATOM	2175	CD2	HIS	A	280	14.711	33.680	25.779	1.00	34.05	C
ATOM	2176	CE1	HIS	A	280	13.838	34.874	27.395	1.00	33.47	C
ATOM	2177	NE2	HIS	A	280	13.545	34.078	26.384	1.00	33.19	N
ATOM	2178	N	ILE	A	281	16.425	30.892	26.723	1.00	33.62	N
ATOM	2179	CA	ILE	A	281	15.699	29.755	26.176	1.00	33.56	C
ATOM	2180	C	ILE	A	281	14.358	29.554	26.806	1.00	33.01	C
ATOM	2181	O	ILE	A	281	14.242	29.475	28.012	1.00	32.55	O
ATOM	2182	CB	ILE	A	281	16.554	28.501	26.300	1.00	33.91	C
ATOM	2183	CG1	ILE	A	281	17.699	28.600	25.295	1.00	35.00	C
ATOM	2184	CG2	ILE	A	281	15.752	27.277	25.988	1.00	34.05	C
ATOM	2185	CD1	ILE	A	281	18.797	27.663	25.562	1.00	36.33	C
ATOM	2186	N	GLU	A	282	13.343	29.445	25.960	1.00	33.12	N
ATOM	2187	CA	GLU	A	282	11.977	29.342	26.441	1.00	33.13	C
ATOM	2188	C	GLU	A	282	11.168	28.219	25.777	1.00	32.96	C
ATOM	2189	O	GLU	A	282	11.252	27.988	24.577	1.00	32.94	O
ATOM	2190	CB	GLU	A	282	11.290	30.707	26.327	1.00	33.06	C
ATOM	2191	CG	GLU	A	282	11.065	31.250	24.925	1.00	33.43	C
ATOM	2192	CD	GLU	A	282	10.529	32.682	24.951	1.00	33.96	C
ATOM	2193	OE1	GLU	A	282	11.071	33.500	25.703	1.00	34.61	O
ATOM	2194	OE2	GLU	A	282	9.563	33.009	24.236	1.00	35.15	O
ATOM	2195	N	SER	A	283	10.431	27.497	26.607	1.00	32.96	N
ATOM	2196	CA	SER	A	283	9.571	26.420	26.169	1.00	33.51	C
ATOM	2197	C	SER	A	283	8.247	27.070	25.784	1.00	34.03	C
ATOM	2198	O	SER	A	283	7.654	27.772	26.597	1.00	34.19	O
ATOM	2199	CB	SER	A	283	9.375	25.408	27.303	1.00	33.43	C
ATOM	2200	OG	SER	A	283	10.371	24.393	27.280	1.00	32.99	O
ATOM	2201	N	LEU	A	284	7.769	26.828	24.565	1.00	34.36	N
ATOM	2202	CA	LEU	A	284	6.610	27.553	24.081	1.00	34.81	C
ATOM	2203	C	LEU	A	284	5.399	27.454	24.982	1.00	35.14	C
ATOM	2204	O	LEU	A	284	5.128	26.425	25.599	1.00	35.08	O
ATOM	2205	CB	LEU	A	284	6.226	27.131	22.670	1.00	35.02	C
ATOM	2206	CG	LEU	A	284	7.302	27.255	21.596	1.00	35.52	C

ATOM	2207	CD1	LEU	A	284	6.675	27.427	20.231	1.00	36.06	C
ATOM	2208	CD2	LEU	A	284	8.197	28.404	21.862	1.00	36.37	C
ATOM	2209	N	LEU	A	285	4.674	28.561	25.040	1.00	35.48	N
ATOM	2210	CA	LEU	A	285	3.481	28.641	25.832	1.00	35.85	C
ATOM	2211	C	LEU	A	285	2.509	27.670	25.255	1.00	36.11	C
ATOM	2212	O	LEU	A	285	2.314	27.623	24.041	1.00	36.34	O
ATOM	2213	CB	LEU	A	285	2.876	30.024	25.742	1.00	35.90	C
ATOM	2214	CG	LEU	A	285	3.740	31.116	26.343	1.00	36.22	C
ATOM	2215	CD1	LEU	A	285	3.271	32.449	25.821	1.00	36.13	C
ATOM	2216	CD2	LEU	A	285	3.701	31.034	27.861	1.00	36.27	C
ATOM	2217	N	ASN	A	286	1.909	26.879	26.128	1.00	36.14	N
ATOM	2218	CA	ASN	A	286	0.890	25.958	25.703	1.00	36.04	C
ATOM	2219	C	ASN	A	286	1.373	24.932	24.663	1.00	35.65	C
ATOM	2220	O	ASN	A	286	0.593	24.471	23.833	1.00	35.84	O
ATOM	2221	CB	ASN	A	286	-0.283	26.792	25.191	1.00	36.33	C
ATOM	2222	CG	ASN	A	286	-0.805	27.766	26.254	1.00	36.97	C
ATOM	2223	OD1	ASN	A	286	-1.266	27.346	27.316	1.00	37.84	O
ATOM	2224	ND2	ASN	A	286	-0.716	29.067	25.977	1.00	37.29	N
ATOM	2225	N	GLY	A	287	2.648	24.550	24.736	1.00	35.18	N
ATOM	2226	CA	GLY	A	287	3.220	23.588	23.806	1.00	34.71	C
ATOM	2227	C	GLY	A	287	3.556	22.252	24.428	1.00	34.44	C
ATOM	2228	O	GLY	A	287	4.071	21.349	23.764	1.00	34.53	O
ATOM	2229	N	GLY	A	288	3.255	22.106	25.706	1.00	34.18	N
ATOM	2230	CA	GLY	A	288	3.521	20.858	26.371	1.00	34.22	C
ATOM	2231	C	GLY	A	288	4.964	20.793	26.766	1.00	34.25	C
ATOM	2232	O	GLY	A	288	5.727	21.719	26.523	1.00	34.28	O
ATOM	2233	N	ILE	A	289	5.351	19.674	27.349	1.00	34.55	N
ATOM	2234	CA	ILE	A	289	6.671	19.572	27.918	1.00	34.83	C
ATOM	2235	C	ILE	A	289	7.702	19.529	26.837	1.00	34.87	C
ATOM	2236	O	ILE	A	289	7.417	19.196	25.687	1.00	35.21	O
ATOM	2237	CB	ILE	A	289	6.815	18.325	28.786	1.00	35.01	C
ATOM	2238	CG1	ILE	A	289	7.018	17.091	27.924	1.00	35.30	C
ATOM	2239	CG2	ILE	A	289	5.610	18.143	29.694	1.00	35.22	C
ATOM	2240	CD1	ILE	A	289	7.654	15.972	28.698	1.00	35.70	C
ATOM	2241	N	THR	A	290	8.921	19.853	27.219	1.00	34.80	N
ATOM	2242	CA	THR	A	290	10.001	19.849	26.273	1.00	34.64	C
ATOM	2243	C	THR	A	290	11.103	18.951	26.740	1.00	34.30	C
ATOM	2244	O	THR	A	290	11.286	18.742	27.924	1.00	34.66	O
ATOM	2245	CB	THR	A	290	10.522	21.258	26.094	1.00	34.75	C
ATOM	2246	OG1	THR	A	290	10.747	21.863	27.375	1.00	34.33	O
ATOM	2247	CG2	THR	A	290	9.455	22.105	25.458	1.00	34.98	C
ATOM	2248	N	ILE	A	291	11.832	18.395	25.802	1.00	33.95	N
ATOM	2249	CA	ILE	A	291	12.940	17.576	26.176	1.00	34.07	C
ATOM	2250	C	ILE	A	291	14.091	17.963	25.310	1.00	33.73	C
ATOM	2251	O	ILE	A	291	13.933	18.203	24.125	1.00	33.83	O
ATOM	2252	CB	ILE	A	291	12.590	16.111	26.029	1.00	34.23	C
ATOM	2253	CG1	ILE	A	291	11.439	15.793	26.967	1.00	34.84	C
ATOM	2254	CG2	ILE	A	291	13.758	15.248	26.429	1.00	34.60	C
ATOM	2255	CD1	ILE	A	291	11.021	14.384	26.946	1.00	35.22	C
ATOM	2256	N	THR	A	292	15.254	18.062	25.913	1.00	33.47	N
ATOM	2257	CA	THR	A	292	16.410	18.418	25.158	1.00	33.68	C
ATOM	2258	C	THR	A	292	17.579	17.741	25.748	1.00	33.73	C
ATOM	2259	O	THR	A	292	17.623	17.472	26.942	1.00	33.28	O
ATOM	2260	CB	THR	A	292	16.725	19.921	25.265	1.00	33.93	C
ATOM	2261	OG1	THR	A	292	15.590	20.725	24.921	1.00	33.80	O
ATOM	2262	CG2	THR	A	292	17.774	20.293	24.245	1.00	34.05	C
ATOM	2263	N	VAL	A	293	18.565	17.518	24.906	1.00	34.13	N
ATOM	2264	CA	VAL	A	293	19.821	17.030	25.391	1.00	34.56	C
ATOM	2265	C	VAL	A	293	20.935	17.867	24.813	1.00	34.79	C
ATOM	2266	O	VAL	A	293	21.039	18.017	23.600	1.00	34.88	O
ATOM	2267	CB	VAL	A	293	20.029	15.603	25.019	1.00	34.66	C

ATOM	2268	CG1	VAL	A	293	21.473	15.238	25.222	1.00	35.24	C
ATOM	2269	CG2	VAL	A	293	19.141	14.741	25.880	1.00	35.15	C
ATOM	2270	N	ASN	A	294	21.768	18.433	25.681	1.00	34.82	N
ATOM	2271	CA	ASN	A	294	22.890	19.187	25.174	1.00	34.61	C
ATOM	2272	C	ASN	A	294	24.159	18.372	25.251	1.00	34.68	C
ATOM	2273	O	ASN	A	294	24.220	17.350	25.934	1.00	34.28	O
ATOM	2274	CB	ASN	A	294	23.036	20.571	25.829	1.00	34.77	C
ATOM	2275	CG	ASN	A	294	23.676	20.533	27.197	1.00	33.90	C
ATOM	2276	OD1	ASN	A	294	24.170	19.504	27.653	1.00	36.35	O
ATOM	2277	ND2	ASN	A	294	23.658	21.670	27.868	1.00	30.46	N
ATOM	2278	N	PHE	A	295	25.141	18.870	24.510	1.00	34.93	N
ATOM	2279	CA	PHE	A	295	26.450	18.291	24.326	1.00	35.27	C
ATOM	2280	C	PHE	A	295	27.467	19.411	24.552	1.00	35.69	C
ATOM	2281	O	PHE	A	295	27.610	20.288	23.698	1.00	35.68	O
ATOM	2282	CB	PHE	A	295	26.581	17.832	22.875	1.00	35.33	C
ATOM	2283	CG	PHE	A	295	25.858	16.554	22.545	1.00	34.85	C
ATOM	2284	CD1	PHE	A	295	24.492	16.549	22.332	1.00	35.36	C
ATOM	2285	CD2	PHE	A	295	26.559	15.371	22.383	1.00	33.46	C
ATOM	2286	CE1	PHE	A	295	23.841	15.378	21.996	1.00	35.19	C
ATOM	2287	CE2	PHE	A	295	25.918	14.207	22.048	1.00	33.20	C
ATOM	2288	CZ	PHE	A	295	24.566	14.202	21.853	1.00	34.09	C
ATOM	2289	N	TRP	A	296	28.182	19.369	25.677	1.00	36.18	N
ATOM	2290	CA	TRP	A	296	29.113	20.433	26.073	1.00	36.34	C
ATOM	2291	C	TRP	A	296	30.534	19.963	25.938	1.00	36.50	C
ATOM	2292	O	TRP	A	296	30.932	18.969	26.563	1.00	36.32	O
ATOM	2293	CB	TRP	A	296	28.913	20.830	27.541	1.00	36.57	C
ATOM	2294	CG	TRP	A	296	27.934	21.912	27.776	1.00	37.38	C
ATOM	2295	CD1	TRP	A	296	26.997	22.361	26.909	1.00	38.36	C
ATOM	2296	CD2	TRP	A	296	27.787	22.693	28.965	1.00	39.42	C
ATOM	2297	NE1	TRP	A	296	26.275	23.384	27.473	1.00	38.20	N
ATOM	2298	CE2	TRP	A	296	26.737	23.605	28.736	1.00	38.86	C
ATOM	2299	CE3	TRP	A	296	28.450	22.730	30.204	1.00	40.74	C
ATOM	2300	CZ2	TRP	A	296	26.327	24.529	29.682	1.00	41.03	C
ATOM	2301	CZ3	TRP	A	296	28.037	23.665	31.160	1.00	40.86	C
ATOM	2302	CH2	TRP	A	296	26.989	24.544	30.891	1.00	41.52	C
ATOM	2303	N	TYR	A	297	31.303	20.719	25.162	1.00	36.60	N
ATOM	2304	CA	TYR	A	297	32.703	20.442	24.923	1.00	36.79	C
ATOM	2305	C	TYR	A	297	33.534	21.597	25.457	1.00	36.96	C
ATOM	2306	O	TYR	A	297	33.117	22.734	25.363	1.00	36.32	O
ATOM	2307	CB	TYR	A	297	32.923	20.291	23.423	1.00	36.93	C
ATOM	2308	CG	TYR	A	297	32.280	19.054	22.868	1.00	37.30	C
ATOM	2309	CD1	TYR	A	297	30.939	19.046	22.523	1.00	37.28	C
ATOM	2310	CD2	TYR	A	297	33.008	17.880	22.716	1.00	36.72	C
ATOM	2311	CE1	TYR	A	297	30.346	17.907	22.039	1.00	37.15	C
ATOM	2312	CE2	TYR	A	297	32.423	16.747	22.226	1.00	35.89	C
ATOM	2313	CZ	TYR	A	297	31.088	16.764	21.893	1.00	36.57	C
ATOM	2314	OH	TYR	A	297	30.467	15.641	21.405	1.00	36.99	O
ATOM	2315	N	LYS	A	298	34.696	21.303	26.039	1.00	37.93	N
ATOM	2316	CA	LYS	A	298	35.615	22.347	26.495	1.00	38.45	C
ATOM	2317	C	LYS	A	298	36.090	22.994	25.236	1.00	38.43	C
ATOM	2318	O	LYS	A	298	36.245	22.320	24.229	1.00	38.55	O
ATOM	2319	CB	LYS	A	298	36.835	21.782	27.233	1.00	38.67	C
ATOM	2320	CG	LYS	A	298	36.634	21.550	28.737	1.00	40.58	C
ATOM	2321	CD	LYS	A	298	37.928	21.093	29.484	1.00	42.63	C
ATOM	2322	CE	LYS	A	298	37.631	20.622	30.951	1.00	44.00	C
ATOM	2323	NZ	LYS	A	298	38.841	20.251	31.812	1.00	43.61	N
ATOM	2324	N	GLY	A	299	36.347	24.286	25.277	1.00	38.82	N
ATOM	2325	CA	GLY	A	299	36.768	24.976	24.080	1.00	39.19	C
ATOM	2326	C	GLY	A	299	38.174	24.582	23.719	1.00	39.60	C
ATOM	2327	O	GLY	A	299	38.859	23.930	24.502	1.00	39.93	O
ATOM	2328	N	ALA	A	300	38.594	24.963	22.520	1.00	40.20	N

ATOM	2329	CA	ALA	A	300	39.963	24.751	22.072	1.00	40.70	C
ATOM	2330	C	ALA	A	300	40.917	25.588	22.929	1.00	41.18	C
ATOM	2331	O	ALA	A	300	40.483	26.417	23.722	1.00	41.25	O
ATOM	2332	CB	ALA	A	300	40.085	25.143	20.616	1.00	40.64	C
ATOM	2333	N	PRO	A	301	42.215	25.408	22.733	1.00	41.91	N
ATOM	2334	CA	PRO	A	301	43.213	26.102	23.542	1.00	42.49	C
ATOM	2335	C	PRO	A	301	43.448	27.541	23.128	1.00	43.14	C
ATOM	2336	O	PRO	A	301	43.323	27.899	21.962	1.00	43.31	O
ATOM	2337	CB	PRO	A	301	44.483	25.326	23.230	1.00	42.38	C
ATOM	2338	CG	PRO	A	301	44.300	24.924	21.833	1.00	42.16	C
ATOM	2339	CD	PRO	A	301	42.851	24.571	21.701	1.00	41.99	C
ATOM	2340	N	THR	A	302	43.823	28.343	24.112	1.00	43.86	N
ATOM	2341	CA	THR	A	302	44.199	29.726	23.915	1.00	44.26	C
ATOM	2342	C	THR	A	302	45.588	29.732	23.318	1.00	44.56	C
ATOM	2343	O	THR	A	302	46.524	29.337	24.011	1.00	44.82	O
ATOM	2344	CB	THR	A	302	44.284	30.383	25.277	1.00	44.20	C
ATOM	2345	OG1	THR	A	302	43.030	30.253	25.956	1.00	45.23	O
ATOM	2346	CG2	THR	A	302	44.512	31.858	25.138	1.00	44.53	C
ATOM	2347	N	PRO	A	303	45.770	30.229	22.091	1.00	44.72	N
ATOM	2348	CA	PRO	A	303	47.083	30.128	21.438	1.00	44.51	C
ATOM	2349	C	PRO	A	303	48.177	30.759	22.306	1.00	44.48	C
ATOM	2350	O	PRO	A	303	47.835	31.522	23.221	1.00	44.02	O
ATOM	2351	CB	PRO	A	303	46.897	30.929	20.145	1.00	44.55	C
ATOM	2352	CG	PRO	A	303	45.426	30.929	19.904	1.00	44.28	C
ATOM	2353	CD	PRO	A	303	44.834	31.040	21.288	1.00	44.71	C
ATOM	2354	N	GLU	A	307	46.829	36.983	18.732	1.00	42.48	N
ATOM	2355	CA	GLU	A	307	46.845	37.952	17.646	1.00	42.70	C
ATOM	2356	C	GLU	A	307	45.812	39.039	17.801	1.00	42.57	C
ATOM	2357	O	GLU	A	307	44.654	38.807	18.138	1.00	42.62	O
ATOM	2358	CB	GLU	A	307	46.624	37.296	16.284	1.00	43.00	C
ATOM	2359	CG	GLU	A	307	46.980	38.235	15.127	1.00	43.85	C
ATOM	2360	CD	GLU	A	307	47.091	37.541	13.779	1.00	44.71	C
ATOM	2361	OE1	GLU	A	307	46.618	36.385	13.643	1.00	45.10	O
ATOM	2362	OE2	GLU	A	307	47.649	38.170	12.851	1.00	45.60	O
ATOM	2363	N	TYR	A	308	46.249	40.243	17.503	1.00	42.66	N
ATOM	2364	CA	TYR	A	308	45.384	41.379	17.577	1.00	42.64	C
ATOM	2365	C	TYR	A	308	44.814	41.530	16.189	1.00	42.85	C
ATOM	2366	O	TYR	A	308	45.413	41.066	15.203	1.00	43.43	O
ATOM	2367	CB	TYR	A	308	46.184	42.565	18.034	1.00	42.47	C
ATOM	2368	CG	TYR	A	308	46.744	42.281	19.399	1.00	43.35	C
ATOM	2369	CD1	TYR	A	308	45.996	42.552	20.534	1.00	44.58	C
ATOM	2370	CD2	TYR	A	308	47.986	41.681	19.559	1.00	43.97	C
ATOM	2371	CE1	TYR	A	308	46.482	42.284	21.790	1.00	45.23	C
ATOM	2372	CE2	TYR	A	308	48.483	41.401	20.818	1.00	44.79	C
ATOM	2373	CZ	TYR	A	308	47.726	41.708	21.935	1.00	45.46	C
ATOM	2374	OH	TYR	A	308	48.200	41.453	23.207	1.00	45.86	O
ATOM	2375	N	PRO	A	309	43.631	42.119	16.111	1.00	42.47	N
ATOM	2376	CA	PRO	A	309	42.930	42.648	17.285	1.00	42.08	C
ATOM	2377	C	PRO	A	309	42.209	41.528	18.051	1.00	41.24	C
ATOM	2378	O	PRO	A	309	41.774	40.578	17.408	1.00	40.92	O
ATOM	2379	CB	PRO	A	309	41.910	43.583	16.652	1.00	42.39	C
ATOM	2380	CG	PRO	A	309	41.574	42.895	15.337	1.00	42.16	C
ATOM	2381	CD	PRO	A	309	42.843	42.260	14.874	1.00	42.28	C
ATOM	2382	N	LEU	A	310	42.059	41.654	19.372	1.00	40.25	N
ATOM	2383	CA	LEU	A	310	41.436	40.596	20.180	1.00	39.46	C
ATOM	2384	C	LEU	A	310	39.971	40.434	19.884	1.00	38.93	C
ATOM	2385	O	LEU	A	310	39.288	41.412	19.624	1.00	39.65	O
ATOM	2386	CB	LEU	A	310	41.526	40.916	21.658	1.00	39.21	C
ATOM	2387	CG	LEU	A	310	42.486	40.099	22.508	1.00	38.87	C
ATOM	2388	CD1	LEU	A	310	43.727	39.738	21.750	1.00	38.58	C
ATOM	2389	CD2	LEU	A	310	42.817	40.926	23.735	1.00	39.01	C

ATOM	2390	N	LYS	A	311	39.462	39.217	19.957	1.00	37.90	N
ATOM	2391	CA	LYS	A	311	38.036	39.029	19.731	1.00	37.53	C
ATOM	2392	C	LYS	A	311	37.217	39.359	20.978	1.00	36.85	C
ATOM	2393	O	LYS	A	311	37.705	39.272	22.092	1.00	36.85	O
ATOM	2394	CB	LYS	A	311	37.746	37.609	19.274	1.00	37.54	C
ATOM	2395	N	ALA	A	312	35.965	39.736	20.793	1.00	36.18	N
ATOM	2396	CA	ALA	A	312	35.112	40.015	21.937	1.00	35.92	C
ATOM	2397	C	ALA	A	312	35.232	38.994	23.058	1.00	35.88	C
ATOM	2398	O	ALA	A	312	35.347	39.384	24.213	1.00	35.99	O
ATOM	2399	CB	ALA	A	312	33.665	40.115	21.526	1.00	35.50	C
ATOM	2400	N	HIS	A	313	35.221	37.700	22.739	1.00	35.71	N
ATOM	2401	CA	HIS	A	313	35.136	36.709	23.797	1.00	35.52	C
ATOM	2402	C	HIS	A	313	36.429	36.694	24.517	1.00	35.31	C
ATOM	2403	O	HIS	A	313	36.489	36.278	25.667	1.00	35.53	O
ATOM	2404	CB	HIS	A	313	34.768	35.291	23.313	1.00	35.89	C
ATOM	2405	CG	HIS	A	313	35.844	34.613	22.535	1.00	36.41	C
ATOM	2406	ND1	HIS	A	313	36.097	34.905	21.214	1.00	38.71	N
ATOM	2407	CD2	HIS	A	313	36.741	33.665	22.889	1.00	37.90	C
ATOM	2408	CE1	HIS	A	313	37.111	34.175	20.789	1.00	38.65	C
ATOM	2409	NE2	HIS	A	313	37.524	33.416	21.788	1.00	38.60	N
ATOM	2410	N	GLN	A	314	37.478	37.162	23.861	1.00	35.08	N
ATOM	2411	CA	GLN	A	314	38.760	37.203	24.535	1.00	35.00	C
ATOM	2412	C	GLN	A	314	38.706	38.301	25.576	1.00	34.62	C
ATOM	2413	O	GLN	A	314	39.145	38.109	26.703	1.00	34.28	O
ATOM	2414	CB	GLN	A	314	39.924	37.399	23.558	1.00	35.10	C
ATOM	2415	CG	GLN	A	314	40.091	36.229	22.597	1.00	35.42	C
ATOM	2416	CD	GLN	A	314	41.107	36.478	21.500	1.00	35.10	C
ATOM	2417	OE1	GLN	A	314	40.938	37.377	20.671	1.00	35.86	O
ATOM	2418	NE2	GLN	A	314	42.150	35.666	21.476	1.00	34.32	N
ATOM	2419	N	LYS	A	315	38.173	39.454	25.197	1.00	34.70	N
ATOM	2420	CA	LYS	A	315	38.009	40.539	26.152	1.00	35.01	C
ATOM	2421	C	LYS	A	315	37.164	40.063	27.320	1.00	34.67	C
ATOM	2422	O	LYS	A	315	37.485	40.325	28.467	1.00	34.81	O
ATOM	2423	CB	LYS	A	315	37.395	41.766	25.501	1.00	35.19	C
ATOM	2424	CG	LYS	A	315	38.344	42.447	24.563	1.00	35.85	C
ATOM	2425	CD	LYS	A	315	37.703	43.636	23.931	1.00	37.25	C
ATOM	2426	CE	LYS	A	315	38.657	44.348	23.001	1.00	39.28	C
ATOM	2427	NZ	LYS	A	315	37.969	45.430	22.222	1.00	40.34	N
ATOM	2428	N	VAL	A	316	36.117	39.314	27.040	1.00	34.30	N
ATOM	2429	CA	VAL	A	316	35.321	38.787	28.118	1.00	34.31	C
ATOM	2430	C	VAL	A	316	36.175	37.925	29.047	1.00	34.37	C
ATOM	2431	O	VAL	A	316	36.071	38.023	30.273	1.00	34.57	O
ATOM	2432	CB	VAL	A	316	34.138	37.992	27.606	1.00	34.39	C
ATOM	2433	CG1	VAL	A	316	33.334	37.434	28.794	1.00	34.91	C
ATOM	2434	CG2	VAL	A	316	33.258	38.873	26.719	1.00	33.60	C
ATOM	2435	N	ALA	A	317	37.011	37.072	28.475	1.00	34.24	N
ATOM	2436	CA	ALA	A	317	37.907	36.267	29.288	1.00	34.17	C
ATOM	2437	C	ALA	A	317	38.744	37.160	30.166	1.00	34.10	C
ATOM	2438	O	ALA	A	317	38.919	36.906	31.352	1.00	33.87	O
ATOM	2439	CB	ALA	A	317	38.813	35.459	28.431	1.00	34.02	C
ATOM	2440	N	ILE	A	318	39.274	38.216	29.574	1.00	34.29	N
ATOM	2441	CA	ILE	A	318	40.178	39.062	30.322	1.00	34.26	C
ATOM	2442	C	ILE	A	318	39.467	39.611	31.546	1.00	33.99	C
ATOM	2443	O	ILE	A	318	39.993	39.556	32.661	1.00	33.96	O
ATOM	2444	CB	ILE	A	318	40.755	40.171	29.447	1.00	33.98	C
ATOM	2445	CG1	ILE	A	318	41.775	39.558	28.489	1.00	34.02	C
ATOM	2446	CG2	ILE	A	318	41.429	41.188	30.323	1.00	33.92	C
ATOM	2447	CD1	ILE	A	318	42.356	40.498	27.428	1.00	33.97	C
ATOM	2448	N	MET	A	319	38.241	40.068	31.337	1.00	33.55	N
ATOM	2449	CA	MET	A	319	37.499	40.741	32.383	1.00	33.41	C
ATOM	2450	C	MET	A	319	37.196	39.772	33.496	1.00	33.38	C

ATOM	2451	O	MET	A	319	37.238	40.101	34.669	1.00	33.32	O
ATOM	2452	CB	MET	A	319	36.193	41.308	31.827	1.00	33.31	C
ATOM	2453	CG	MET	A	319	36.361	42.521	30.951	1.00	32.96	C
ATOM	2454	SD	MET	A	319	34.812	43.355	30.617	1.00	33.05	S
ATOM	2455	CE	MET	A	319	34.119	42.299	29.410	1.00	33.43	C
ATOM	2456	N	ARG	A	320	36.873	38.558	33.121	1.00	33.68	N
ATOM	2457	CA	ARG	A	320	36.575	37.568	34.112	1.00	33.61	C
ATOM	2458	C	ARG	A	320	37.796	37.392	34.996	1.00	33.82	C
ATOM	2459	O	ARG	A	320	37.719	37.448	36.225	1.00	33.48	O
ATOM	2460	CB	ARG	A	320	36.211	36.292	33.402	1.00	33.39	C
ATOM	2461	CG	ARG	A	320	34.875	36.406	32.725	1.00	33.88	C
ATOM	2462	CD	ARG	A	320	34.268	35.080	32.340	1.00	34.20	C
ATOM	2463	NE	ARG	A	320	34.217	34.215	33.507	1.00	33.14	N
ATOM	2464	CZ	ARG	A	320	33.159	34.103	34.260	1.00	33.15	C
ATOM	2465	NH1	ARG	A	320	32.077	34.762	33.933	1.00	33.83	N
ATOM	2466	NH2	ARG	A	320	33.170	33.338	35.331	1.00	34.46	N
ATOM	2467	N	ASN	A	321	38.941	37.232	34.355	1.00	33.92	N
ATOM	2468	CA	ASN	A	321	40.143	36.978	35.085	1.00	34.08	C
ATOM	2469	C	ASN	A	321	40.486	38.141	36.018	1.00	34.26	C
ATOM	2470	O	ASN	A	321	40.874	37.919	37.164	1.00	34.14	O
ATOM	2471	CB	ASN	A	321	41.271	36.616	34.119	1.00	34.30	C
ATOM	2472	CG	ASN	A	321	41.195	35.161	33.663	1.00	34.44	C
ATOM	2473	OD1	ASN	A	321	40.829	34.273	34.434	1.00	34.16	O
ATOM	2474	ND2	ASN	A	321	41.522	34.918	32.402	1.00	35.05	N
ATOM	2475	N	ILE	A	322	40.318	39.379	35.569	1.00	34.24	N
ATOM	2476	CA	ILE	A	322	40.634	40.488	36.448	1.00	34.25	C
ATOM	2477	C	ILE	A	322	39.781	40.408	37.701	1.00	33.85	C
ATOM	2478	O	ILE	A	322	40.250	40.711	38.789	1.00	34.16	O
ATOM	2479	CB	ILE	A	322	40.412	41.837	35.760	1.00	34.70	C
ATOM	2480	CG1	ILE	A	322	41.338	41.993	34.558	1.00	35.34	C
ATOM	2481	CG2	ILE	A	322	40.651	43.007	36.748	1.00	35.37	C
ATOM	2482	CD1	ILE	A	322	42.778	41.928	34.895	1.00	36.17	C
ATOM	2483	N	GLU	A	323	38.525	40.013	37.558	1.00	33.42	N
ATOM	2484	CA	GLU	A	323	37.636	39.935	38.704	1.00	32.97	C
ATOM	2485	C	GLU	A	323	38.085	38.842	39.652	1.00	33.11	C
ATOM	2486	O	GLU	A	323	38.067	39.001	40.864	1.00	32.66	O
ATOM	2487	CB	GLU	A	323	36.199	39.673	38.256	1.00	32.73	C
ATOM	2488	CG	GLU	A	323	35.543	40.907	37.672	1.00	33.02	C
ATOM	2489	CD	GLU	A	323	34.166	40.693	37.057	1.00	32.91	C
ATOM	2490	OE1	GLU	A	323	33.193	40.309	37.740	1.00	31.94	O
ATOM	2491	OE2	GLU	A	323	34.048	40.961	35.856	1.00	33.81	O
ATOM	2492	N	LYS	A	324	38.497	37.720	39.088	1.00	33.56	N
ATOM	2493	CA	LYS	A	324	38.882	36.588	39.895	1.00	33.72	C
ATOM	2494	C	LYS	A	324	40.138	36.937	40.670	1.00	34.09	C
ATOM	2495	O	LYS	A	324	40.228	36.664	41.859	1.00	34.15	O
ATOM	2496	CB	LYS	A	324	39.119	35.368	39.012	1.00	33.73	C
ATOM	2497	CG	LYS	A	324	37.877	34.766	38.353	1.00	33.83	C
ATOM	2498	CD	LYS	A	324	38.355	33.673	37.440	1.00	33.70	C
ATOM	2499	CE	LYS	A	324	37.294	32.883	36.723	1.00	33.57	C
ATOM	2500	NZ	LYS	A	324	37.993	31.761	35.984	1.00	32.76	N
ATOM	2501	N	MET	A	325	41.098	37.558	39.995	1.00	34.73	N
ATOM	2502	CA	MET	A	325	42.380	37.915	40.609	1.00	35.32	C
ATOM	2503	C	MET	A	325	42.248	38.883	41.766	1.00	35.06	C
ATOM	2504	O	MET	A	325	42.888	38.729	42.808	1.00	34.50	O
ATOM	2505	CB	MET	A	325	43.328	38.501	39.563	1.00	35.57	C
ATOM	2506	CG	MET	A	325	43.962	37.419	38.724	1.00	36.96	C
ATOM	2507	SD	MET	A	325	44.984	37.973	37.360	1.00	40.43	S
ATOM	2508	CE	MET	A	325	45.710	39.363	38.037	1.00	41.05	C
ATOM	2509	N	LEU	A	326	41.427	39.891	41.555	1.00	35.32	N
ATOM	2510	CA	LEU	A	326	41.178	40.895	42.570	1.00	35.83	C
ATOM	2511	C	LEU	A	326	40.556	40.230	43.770	1.00	35.60	C

ATOM	2512	O	LEU	A	326	40.914	40.507	44.910	1.00	35.34	O
ATOM	2513	CB	LEU	A	326	40.216	41.929	42.026	1.00	36.07	C
ATOM	2514	CG	LEU	A	326	40.721	43.321	41.689	1.00	37.48	C
ATOM	2515	CD1	LEU	A	326	42.221	43.442	41.537	1.00	38.59	C
ATOM	2516	CD2	LEU	A	326	40.049	43.692	40.413	1.00	38.65	C
ATOM	2517	N	GLY	A	327	39.613	39.341	43.493	1.00	35.59	N
ATOM	2518	CA	GLY	A	327	38.947	38.604	44.538	1.00	35.54	C
ATOM	2519	C	GLY	A	327	39.983	37.998	45.443	1.00	35.62	C
ATOM	2520	O	GLY	A	327	40.023	38.258	46.641	1.00	35.52	O
ATOM	2521	N	GLU	A	328	40.863	37.209	44.851	1.00	35.84	N
ATOM	2522	CA	GLU	A	328	41.831	36.486	45.632	1.00	35.91	C
ATOM	2523	C	GLU	A	328	42.783	37.424	46.298	1.00	35.44	C
ATOM	2524	O	GLU	A	328	43.086	37.272	47.465	1.00	35.60	O
ATOM	2525	CB	GLU	A	328	42.584	35.513	44.750	1.00	36.21	C
ATOM	2526	CG	GLU	A	328	41.675	34.419	44.226	1.00	38.20	C
ATOM	2527	CD	GLU	A	328	41.854	33.088	44.941	1.00	41.33	C
ATOM	2528	OE1	GLU	A	328	42.244	33.084	46.138	1.00	42.40	O
ATOM	2529	OE2	GLU	A	328	41.608	32.034	44.292	1.00	43.59	O
ATOM	2530	N	ALA	A	329	43.220	38.435	45.577	1.00	35.11	N
ATOM	2531	CA	ALA	A	329	44.259	39.282	46.110	1.00	34.85	C
ATOM	2532	C	ALA	A	329	43.790	40.103	47.296	1.00	34.75	C
ATOM	2533	O	ALA	A	329	44.533	40.282	48.258	1.00	34.53	O
ATOM	2534	CB	ALA	A	329	44.778	40.163	45.046	1.00	34.85	C
ATOM	2535	N	LEU	A	330	42.567	40.613	47.214	1.00	34.62	N
ATOM	2536	CA	LEU	A	330	41.996	41.399	48.291	1.00	34.55	C
ATOM	2537	C	LEU	A	330	41.478	40.555	49.448	1.00	34.90	C
ATOM	2538	O	LEU	A	330	41.234	41.072	50.532	1.00	35.33	O
ATOM	2539	CB	LEU	A	330	40.851	42.245	47.766	1.00	34.31	C
ATOM	2540	CG	LEU	A	330	41.202	43.322	46.743	1.00	34.06	C
ATOM	2541	CD1	LEU	A	330	39.942	43.965	46.258	1.00	34.23	C
ATOM	2542	CD2	LEU	A	330	42.078	44.382	47.285	1.00	34.05	C
ATOM	2543	N	GLY	A	331	41.295	39.262	49.232	1.00	35.23	N
ATOM	2544	CA	GLY	A	331	40.794	38.401	50.285	1.00	35.49	C
ATOM	2545	C	GLY	A	331	39.306	38.471	50.584	1.00	35.66	C
ATOM	2546	O	GLY	A	331	38.849	37.859	51.544	1.00	35.61	O
ATOM	2547	N	ASN	A	332	38.547	39.213	49.790	1.00	36.04	N
ATOM	2548	CA	ASN	A	332	37.113	39.295	49.992	1.00	36.44	C
ATOM	2549	C	ASN	A	332	36.483	39.791	48.732	1.00	36.55	C
ATOM	2550	O	ASN	A	332	36.647	40.951	48.381	1.00	37.01	O
ATOM	2551	CB	ASN	A	332	36.775	40.275	51.094	1.00	36.54	C
ATOM	2552	CG	ASN	A	332	35.311	40.260	51.440	1.00	37.47	C
ATOM	2553	OD1	ASN	A	332	34.519	39.541	50.828	1.00	37.83	O
ATOM	2554	ND2	ASN	A	332	34.938	41.043	52.444	1.00	39.90	N
ATOM	2555	N	PRO	A	333	35.699	38.957	48.078	1.00	36.46	N
ATOM	2556	CA	PRO	A	333	35.178	39.338	46.775	1.00	36.43	C
ATOM	2557	C	PRO	A	333	34.368	40.590	46.908	1.00	36.32	C
ATOM	2558	O	PRO	A	333	34.318	41.384	45.988	1.00	36.36	O
ATOM	2559	CB	PRO	A	333	34.278	38.170	46.366	1.00	36.50	C
ATOM	2560	CG	PRO	A	333	34.247	37.229	47.489	1.00	36.58	C
ATOM	2561	CD	PRO	A	333	35.180	37.668	48.548	1.00	36.54	C
ATOM	2562	N	GLN	A	334	33.743	40.800	48.047	1.00	36.44	N
ATOM	2563	CA	GLN	A	334	32.906	41.969	48.130	1.00	36.85	C
ATOM	2564	C	GLN	A	334	33.748	43.222	48.057	1.00	36.57	C
ATOM	2565	O	GLN	A	334	33.218	44.304	47.868	1.00	36.98	O
ATOM	2566	CB	GLN	A	334	31.987	41.940	49.348	1.00	37.31	C
ATOM	2567	CG	GLN	A	334	30.685	41.151	49.026	1.00	38.95	C
ATOM	2568	CD	GLN	A	334	29.669	41.097	50.171	1.00	40.33	C
ATOM	2569	OE1	GLN	A	334	29.751	41.872	51.137	1.00	41.70	O
ATOM	2570	NE2	GLN	A	334	28.712	40.178	50.061	1.00	39.80	N
ATOM	2571	N	GLU	A	335	35.064	43.088	48.159	1.00	35.99	N
ATOM	2572	CA	GLU	A	335	35.914	44.261	48.067	1.00	35.53	C

ATOM	2573	C	GLU	A	335	36.229	44.610	46.621	1.00	34.53	C
ATOM	2574	O	GLU	A	335	36.715	45.689	46.338	1.00	34.20	O
ATOM	2575	CB	GLU	A	335	37.201	44.083	48.894	1.00	35.99	C
ATOM	2576	CG	GLU	A	335	37.054	44.507	50.354	1.00	37.38	C
ATOM	2577	CD	GLU	A	335	38.382	44.592	51.076	1.00	40.16	C
ATOM	2578	OE1	GLU	A	335	39.271	45.318	50.570	1.00	42.04	O
ATOM	2579	OE2	GLU	A	335	38.535	43.952	52.153	1.00	41.52	O
ATOM	2580	N	VAL	A	336	35.922	43.701	45.706	1.00	33.97	N
ATOM	2581	CA	VAL	A	336	36.165	43.901	44.276	1.00	33.49	C
ATOM	2582	C	VAL	A	336	35.537	45.159	43.670	1.00	33.00	C
ATOM	2583	O	VAL	A	336	36.194	45.904	42.946	1.00	33.01	O
ATOM	2584	CB	VAL	A	336	35.696	42.678	43.473	1.00	33.36	C
ATOM	2585	CG1	VAL	A	336	35.726	42.951	41.992	1.00	34.05	C
ATOM	2586	CG2	VAL	A	336	36.582	41.509	43.755	1.00	33.01	C
ATOM	2587	N	GLY	A	337	34.270	45.406	43.926	1.00	32.41	N
ATOM	2588	CA	GLY	A	337	33.675	46.592	43.351	1.00	32.21	C
ATOM	2589	C	GLY	A	337	34.332	47.903	43.724	1.00	31.72	C
ATOM	2590	O	GLY	A	337	34.682	48.711	42.859	1.00	31.40	O
ATOM	2591	N	PRO	A	338	34.433	48.163	45.013	1.00	31.50	N
ATOM	2592	CA	PRO	A	338	35.025	49.415	45.470	1.00	31.47	C
ATOM	2593	C	PRO	A	338	36.387	49.648	44.846	1.00	31.60	C
ATOM	2594	O	PRO	A	338	36.655	50.756	44.386	1.00	31.51	O
ATOM	2595	CB	PRO	A	338	35.065	49.230	46.977	1.00	31.27	C
ATOM	2596	CG	PRO	A	338	33.883	48.376	47.218	1.00	31.08	C
ATOM	2597	CD	PRO	A	338	33.933	47.359	46.137	1.00	31.28	C
ATOM	2598	N	LEU	A	339	37.219	48.620	44.765	1.00	31.80	N
ATOM	2599	CA	LEU	A	339	38.531	48.825	44.182	1.00	32.25	C
ATOM	2600	C	LEU	A	339	38.431	49.269	42.740	1.00	32.12	C
ATOM	2601	O	LEU	A	339	39.130	50.188	42.320	1.00	32.25	O
ATOM	2602	CB	LEU	A	339	39.395	47.581	44.278	1.00	32.49	C
ATOM	2603	CG	LEU	A	339	40.853	47.920	43.962	1.00	33.78	C
ATOM	2604	CD1	LEU	A	339	41.793	47.121	44.797	1.00	34.64	C
ATOM	2605	CD2	LEU	A	339	41.146	47.681	42.516	1.00	34.50	C
ATOM	2606	N	LEU	A	340	37.567	48.607	41.978	1.00	32.12	N
ATOM	2607	CA	LEU	A	340	37.385	48.963	40.584	1.00	31.85	C
ATOM	2608	C	LEU	A	340	36.854	50.376	40.483	1.00	31.56	C
ATOM	2609	O	LEU	A	340	37.342	51.169	39.701	1.00	31.42	O
ATOM	2610	CB	LEU	A	340	36.468	47.975	39.888	1.00	31.90	C
ATOM	2611	CG	LEU	A	340	37.089	46.598	39.669	1.00	32.44	C
ATOM	2612	CD1	LEU	A	340	36.028	45.636	39.132	1.00	33.11	C
ATOM	2613	CD2	LEU	A	340	38.281	46.660	38.728	1.00	31.95	C
ATOM	2614	N	ASN	A	341	35.865	50.717	41.282	1.00	31.61	N
ATOM	2615	CA	ASN	A	341	35.384	52.083	41.231	1.00	31.82	C
ATOM	2616	C	ASN	A	341	36.484	53.109	41.525	1.00	31.45	C
ATOM	2617	O	ASN	A	341	36.600	54.119	40.845	1.00	30.80	O
ATOM	2618	CB	ASN	A	341	34.212	52.260	42.177	1.00	31.97	C
ATOM	2619	CG	ASN	A	341	32.901	51.883	41.532	1.00	32.77	C
ATOM	2620	OD1	ASN	A	341	32.426	52.568	40.614	1.00	34.75	O
ATOM	2621	ND2	ASN	A	341	32.310	50.789	41.992	1.00	31.86	N
ATOM	2622	N	THR	A	342	37.301	52.835	42.530	1.00	31.40	N
ATOM	2623	CA	THR	A	342	38.380	53.739	42.880	1.00	31.27	C
ATOM	2624	C	THR	A	342	39.317	53.842	41.707	1.00	31.14	C
ATOM	2625	O	THR	A	342	39.837	54.908	41.402	1.00	30.66	O
ATOM	2626	CB	THR	A	342	39.146	53.217	44.097	1.00	31.43	C
ATOM	2627	OG1	THR	A	342	38.448	53.565	45.299	1.00	30.95	O
ATOM	2628	CG2	THR	A	342	40.500	53.913	44.228	1.00	31.17	C
ATOM	2629	N	MET	A	343	39.549	52.714	41.059	1.00	31.29	N
ATOM	2630	CA	MET	A	343	40.406	52.703	39.898	1.00	31.59	C
ATOM	2631	C	MET	A	343	39.970	53.639	38.785	1.00	31.49	C
ATOM	2632	O	MET	A	343	40.804	54.361	38.255	1.00	31.46	O
ATOM	2633	CB	MET	A	343	40.516	51.310	39.314	1.00	31.79	C

ATOM	2634	CG	MET	A	343	41.831	50.662	39.629	1.00	32.65	C
ATOM	2635	SD	MET	A	343	42.271	49.379	38.468	1.00	33.09	S
ATOM	2636	CE	MET	A	343	41.116	48.304	38.916	1.00	34.22	C
ATOM	2637	N	ILE	A	344	38.690	53.641	38.418	1.00	31.31	N
ATOM	2638	CA	ILE	A	344	38.293	54.432	37.265	1.00	31.42	C
ATOM	2639	C	ILE	A	344	37.708	55.787	37.533	1.00	31.14	C
ATOM	2640	O	ILE	A	344	37.752	56.630	36.640	1.00	31.01	O
ATOM	2641	CB	ILE	A	344	37.272	53.708	36.362	1.00	31.86	C
ATOM	2642	CG1	ILE	A	344	35.870	53.866	36.928	1.00	33.16	C
ATOM	2643	CG2	ILE	A	344	37.633	52.229	36.118	1.00	32.09	C
ATOM	2644	CD1	ILE	A	344	34.876	53.232	36.072	1.00	35.18	C
ATOM	2645	N	LYS	A	345	37.131	56.050	38.698	1.00	30.88	N
ATOM	2646	CA	LYS	A	345	36.455	57.334	38.758	1.00	30.94	C
ATOM	2647	C	LYS	A	345	37.412	58.468	38.853	1.00	30.72	C
ATOM	2648	O	LYS	A	345	38.250	58.558	39.743	1.00	30.61	O
ATOM	2649	CB	LYS	A	345	35.315	57.475	39.765	1.00	31.14	C
ATOM	2650	CG	LYS	A	345	35.361	56.680	40.977	1.00	32.16	C
ATOM	2651	CD	LYS	A	345	33.970	56.273	41.340	1.00	33.62	C
ATOM	2652	CE	LYS	A	345	33.514	56.873	42.665	1.00	34.71	C
ATOM	2653	NZ	LYS	A	345	34.302	56.317	43.840	1.00	36.28	N
ATOM	2654	N	GLY	A	346	37.280	59.332	37.870	1.00	30.83	N
ATOM	2655	CA	GLY	A	346	38.114	60.504	37.789	1.00	30.66	C
ATOM	2656	C	GLY	A	346	39.345	60.195	37.001	1.00	30.25	C
ATOM	2657	O	GLY	A	346	40.147	61.066	36.745	1.00	30.00	O
ATOM	2658	N	ARG	A	347	39.454	58.953	36.564	1.00	30.42	N
ATOM	2659	CA	ARG	A	347	40.652	58.496	35.923	1.00	30.66	C
ATOM	2660	C	ARG	A	347	40.389	57.879	34.566	1.00	30.89	C
ATOM	2661	O	ARG	A	347	41.033	58.241	33.588	1.00	30.30	O
ATOM	2662	CB	ARG	A	347	41.350	57.526	36.860	1.00	30.88	C
ATOM	2663	CG	ARG	A	347	41.786	58.177	38.171	1.00	30.97	C
ATOM	2664	CD	ARG	A	347	43.129	57.701	38.636	1.00	31.35	C
ATOM	2665	NE	ARG	A	347	43.128	56.255	38.597	1.00	30.88	N
ATOM	2666	CZ	ARG	A	347	44.088	55.482	38.128	1.00	29.55	C
ATOM	2667	NH1	ARG	A	347	45.237	55.962	37.667	1.00	28.44	N
ATOM	2668	NH2	ARG	A	347	43.876	54.181	38.147	1.00	29.95	N
ATOM	2669	N	TYR	A	348	39.446	56.953	34.480	1.00	31.67	N
ATOM	2670	CA	TYR	A	348	39.143	56.377	33.176	1.00	32.27	C
ATOM	2671	C	TYR	A	348	37.726	56.574	32.689	1.00	33.10	C
ATOM	2672	O	TYR	A	348	37.425	56.212	31.557	1.00	33.12	O
ATOM	2673	CB	TYR	A	348	39.422	54.896	33.154	1.00	31.96	C
ATOM	2674	CG	TYR	A	348	40.860	54.527	33.253	1.00	32.06	C
ATOM	2675	CD1	TYR	A	348	41.652	54.424	32.114	1.00	31.60	C
ATOM	2676	CD2	TYR	A	348	41.435	54.256	34.483	1.00	31.75	C
ATOM	2677	CE1	TYR	A	348	42.980	54.061	32.203	1.00	31.08	C
ATOM	2678	CE2	TYR	A	348	42.761	53.897	34.580	1.00	32.11	C
ATOM	2679	CZ	TYR	A	348	43.530	53.800	33.438	1.00	31.05	C
ATOM	2680	OH	TYR	A	348	44.845	53.427	33.544	1.00	29.83	O
ATOM	2681	N	ASN	A	349	36.850	57.137	33.507	1.00	34.48	N
ATOM	2682	CA	ASN	A	349	35.465	57.275	33.083	1.00	35.73	C
ATOM	2683	C	ASN	A	349	35.037	58.643	32.595	1.00	36.60	C
ATOM	2684	O	ASN	A	349	35.757	59.640	32.508	1.00	37.31	O
ATOM	2685	CB	ASN	A	349	34.542	56.900	34.212	1.00	35.62	C
ATOM	2686	CG	ASN	A	349	34.635	57.864	35.353	1.00	37.08	C
ATOM	2687	OD1	ASN	A	349	35.398	58.840	35.308	1.00	36.55	O
ATOM	2688	ND2	ASN	A	349	33.861	57.604	36.403	1.00	39.78	N
ATOM	2689	OXT	ASN	A	349	33.861	58.757	32.276	1.00	38.11	O
TER	2690		ASN	A	349						
HETATM	2691	FE	FE2	A1350		23.364	27.586	28.889	1.00	32.45	FE
HETATM	2692	C1	AKG	A1351		22.523	25.412	27.792	1.00	37.04	C
HETATM	2693	O1	AKG	A1351		23.535	26.092	27.914	1.00	39.03	O
HETATM	2694	O2	AKG	A1351		22.557	24.216	27.226	1.00	37.33	O

HETATM	2695	C2	AKG	A1351	21.186	25.964	28.297	1.00	35.25
HETATM	2696	O5	AKG	A1351	21.175	27.063	28.807	1.00	33.44
HETATM	2697	C3	AKG	A1351	19.937	25.139	28.167	1.00	36.11
HETATM	2698	C4	AKG	A1351	18.848	25.738	29.055	1.00	36.50
HETATM	2699	C5	AKG	A1351	17.464	25.331	28.632	1.00	35.97
HETATM	2700	O3	AKG	A1351	17.276	24.627	27.656	1.00	36.64
HETATM	2701	O4	AKG	A1351	16.432	25.771	29.338	1.00	36.31
HETATM	2702	S	SO4	A1352	0.300	25.196	43.477	1.00	80.75
HETATM	2703	O1	SO4	A1352	1.094	26.117	44.306	1.00	78.50
HETATM	2704	O2	SO4	A1352	1.221	24.413	42.644	1.00	78.51
HETATM	2705	O3	SO4	A1352	-0.470	24.282	44.334	1.00	79.54
HETATM	2706	O4	SO4	A1352	-0.640	25.951	42.633	1.00	78.07
HETATM	2707	S	SO4	A1353	1.825	28.603	29.515	1.00	77.38
HETATM	2708	O1	SO4	A1353	3.041	29.212	30.061	1.00	76.05
HETATM	2709	O2	SO4	A1353	2.144	27.528	28.574	1.00	75.24
HETATM	2710	O3	SO4	A1353	1.090	28.086	30.675	1.00	76.52
HETATM	2711	O4	SO4	A1353	1.011	29.586	28.783	1.00	76.04
HETATM	2712	S	SO4	A1354	34.403	30.284	38.043	1.00	78.17
HETATM	2713	O1	SO4	A1354	34.921	31.567	38.516	1.00	75.93
HETATM	2714	O2	SO4	A1354	35.240	29.158	38.450	1.00	77.42
HETATM	2715	O3	SO4	A1354	33.112	30.084	38.686	1.00	75.59
HETATM	2716	O4	SO4	A1354	34.303	30.307	36.581	1.00	74.60
HETATM	2717	O	HOH	Z 1	25.824	26.442	26.175	1.00	46.43
HETATM	2718	O	HOH	Z 2	27.811	27.709	24.637	1.00	41.32
HETATM	2719	O	HOH	Z 3	27.367	30.233	25.742	1.00	38.18
CONECT	1478	2691							
CONECT	1498	2691							
CONECT	2167	2691							
CONECT	2691	2696	2693	1478	2167	1498			
CONECT	2692	2693	2694	2695					
CONECT	2693	2692	2691						
CONECT	2694	2692							
CONECT	2695	2692	2696	2697					
CONECT	2696	2691	2695						
CONECT	2697	2695	2698						
CONECT	2698	2697	2699						
CONECT	2699	2698	2700	2701					
CONECT	2700	2699							
CONECT	2701	2699							
CONECT	2702	2703	2704	2705	2706				
CONECT	2703	2702							
CONECT	2704	2702							
CONECT	2705	2702							
CONECT	2706	2702							
CONECT	2707	2708	2709	2710	2711				
CONECT	2708	2707							
CONECT	2709	2707							
CONECT	2710	2707							
CONECT	2711	2707							
CONECT	2712	2713	2714	2715	2716				
CONECT	2713	2712							
CONECT	2714	2712							
CONECT	2715	2712							